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Title:
Perfect score:
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MODEL-frame-_n2p.model -DEV-xlp

-MODEL-frame-_n2p.model -DEV-xlp

-O-/cgn2_1/USPTO_Spool/US09778187/runat_20112002_073630_26329/app_guery.fasta_1.2318

-DB-PIR_73 -OPMT-fastan -SUFFIX=rpr -MIRMATCH-0.1 -LOOPCL-0 -LOOPEXT-0

-UNITS-blits -STARP-1 -END--1 -MATRIX-blosum62 -TRANS-human40.cd1 -LIST-45

-DCCALIGN-200 -THR_SCORE-pct -THR_MAX=100 -THR_MIN-0 -ALIGN-15 -MODE-LOCAL

-OUTFMT-pto -NORM-ext -HEAPSIZE=500 -MINLEN-0 -MAXLEN-2000000000

-USER-US09778187_8CGN_1_1_113_8crunat_20112002_073630_26329 -NCPU-6 -ICPU-3

-NO_MAAP -LARGEQUERY -NEG_SCORES-0 -WAIT -LONGLOG -DEV_TIMEOUT=120

-WARN_TIMEOUT=30 -THREADS=1 -XGAPOD=10 -XGAPEXT-0.5 -FGAPEXT-7

-YGAPPOP-10 -YGAPEXT-0.5 -DELOP-6 -DELEXT-7
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Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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### ALIGNMENTS

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R:Kershaw, J.
submitted to the EMBL Data Library,
submitted to umber: Z19929
                   Percent Similarity:
Best Local Similarity:
Query Match:
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A; Introns: 85/1; 120/1; 334/3; 370/1; 477/2; 606/3; 664/1; 935/3; 977/1; 1051/3; 1184
A; Introns: 85/1; 2693/3; 2759/1; 2852/1; 2889/3; 2913/3; 2941/1; 2967/3; 2991/3; 303
1; 4225/1; 4361/1; 4408/1; 4456/1; 4498/1; 4647/3; 4838/1; 4879/1; 4941/1; 5011/1; 50
                                                                                                                                                                                                                                                                                      C;Genetics:
A;Gene: CESP:F15G9.4a
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A;Restidues: 1-5175 <M12>
A;Cross-references: EMBL:247070; PIDN:CAA87344.1; GSPDB:GN00028; CESP:F15G9.
A;Experimental source: clone T09B9
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A; Accession: T24733
A; Status: preliminar
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A;Molecule type: DNA
A;Residues: 1-5175 <WIL>
A;Cross-references: EMBL:247068; PIDN:CAA87335.1; GSPDB:GN00028; CESP:F15G9.4a
A;Experimental source: clone F15G9
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ACCATCCTTACCATCATCACAGATTCCCGAGCAGGTGAA
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| GluThrAspSerSerLysTyrThrCysIleAlaThrAsnGluAlaGlyThrAspLysArg
                                           IleThrValAsnSerGlyAsnProSerThrLeuHisCysProAlaLysGlySerProSer
                                                                                                                                 GATTATATGCTGTATGTATACGATCCCCCCACAACTATCCCTCCT----
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submitted to the EMBL Data
A; Reference number: Z19929
A; Accession: T24734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C; Accessi
R; Vogel,
Qy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-5198 <WIL>
A;Cross-references: EMBL:Z47068; PIDN:CAA87336.1; GSPDB:GN00028; CESP:F15G9.4b
A;Experimental source: clone F15G9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-778-187B-1_COPY_130_1137 (1-1008) x T43290
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DB:
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A; Residues: 1-5198 <WI2>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Percent Similarity:
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A;Experimental source: clone T09B9
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A; Residues: 1-5198 < VOG>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Accession: T43290
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Db
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Gene: him-4; F15G9.4b
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Status: preliminary; translated
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   CTGATGATCGATATCCAGAAAGAC---
                                                                  As n As p Ala Gly Glu As n Ser Phe As p Phe Lys Ala Thr Val Leu Val Pro Pro Thr I leu Val Pro Pro Thr I leu Val Pro Pro Through Control Pro Thro
                                                                                                           ACCGATCCCCCACAGGAAAGTTACACC---ACCATCACAGTCCTGGTCCCACCACGTAAT
::: |||::
                                                                                                                                                                                                      ArgLeuThrIleLeuAsnValThrGluAsnAspGluGlyGlnTyrSerCysArgValLys
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                                                                                                                                                                                                                                                                    IleAspLeuGluAlaGluAspAlaArgIleThrArgLeu-----SerAsnAspArg
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Mismatches:
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2967/3; 2991/3; 303
; 4964/1; 5034/1; 51
      381
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5.1.3

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Command line parameters:

MODEL-framet_n2p.model -DEV-xlp

-Q-/cgn2_1/USPTO_spool/US09778187/runat_20112002_073629_26309/app_query.fasta_1.2318
-Q-/cgn2_1/USPTO_spool/US09778187/runat_20112002_073629_26309/app_query.fasta_1.2318
-DB-SwissProt_40 -QFMT-fastan -SUFFIX-sp -MIMMATCH-0.1 -LOOPCL-0 -LOOPEXT-0

-UNITS-bits -START-1 -END-1 -MATRIX-blosum62 -TRANS-human40.cdi -LIST-45
-DOCALIGN-200 -THR_SCORE-pct -THR_MIN-0 -ALIGN-15 -MODE-LOCAL
-OUTFMT-pto -NORM-ext -HEAPSIZE-50 -MINLEN-0 -MAXIEN-2000000000

-USER-US09778187 eCGN _1_1_45 erunat_20112002_073629_26309 -NCPU-6 -ICPU-3

-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES-0 -WAIT -LONGLOG -DEV_TIMEOUT=120

-WARN_TIMEOUT=30 -THREADS-1 -XGAPOE-10 -XGAPEXT-0.5 -FGAPOP-6 -FGAPEXT-7
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Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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Listing
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                                                                                                                                                                                                                                                                 NCM2_MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -DELEXT-7
                                               OPCM_HUMAN
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does of the state	Q13449	P34082	Q13740	P13596	P31836	08180	P28685	P43121	P15364	P70211	P42292	060500	P13591	P13592	Q98919	Q61490	Q9r044		Q05793	Q90610	P97603	P35329	P32736	Q92859	P06731	P20273	Q90773
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# ALIGNMENTS

\_XENLA NCA1\_XENLA P16170; This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no wav -i- SUBCELLULAR LOCATION: Type I membrane protein.
-i- ALTERNATIVE PRODUCTS: 2 isoforms; N-CAM 180 (sho
-N-CAM 140; are produced by alternative splicing.
-i- TISSUE SPECIFICITY: EXPRESSED IN NEURON AND IN F
TISSUE. Nucleic Acids Res. 17:10321-10335(1989).

Nucleic Acids Res. 17:10321-10335(1989).

-i- FUNCTION: THIS PROTEIN IS A CELL ADHESION MOLECULE INVOLVED

-i- FUNCTION: THIS PROTEIN NEGRITHE FASCICULATION, OUTGROWTH OF SEQUENCE FROM N.A.
MEDLINE-90098871; PubMed-2481269;
Krieg P.A., Sakaguchi D.S., Kintner C.R.;
"Primary structure and developmental expression of a large cytoplasmic domain form of Xenopus laevis neural cell adhesion Xenopus laevis (African clawed frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
Xenopodinae; Xenopus. 01-APR-1990 (Rel. 14, Created)
01-APR-1990 (Rel. 14, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Neural cell adhesion molecule 1, 180 kDa isoform precursor (N-CAM 180). molecule (NCAM)."; NCAM1. NCBI\_TaxID-8355; EARLY NEURAL DEVELOPMENT.

SIMILARITY: BELONGS TO THE IMMUNOGLOBULIN SUPERFAMILY.

SIMILARITY: CONTAINS 5 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAIN

SIMILARITY: CONTAINS 2 FIBRONECTIN TYPE III-LIKE DOMAINS. DEVELOPMENTAL STAGE: THE MRNA ENCODING THIS LD-NCAM IS TRANSCRIPT PRESENT IN BOTH MATERNAL RNA AND IN THE EMBI NEURON-NEURON ADHESION, NEURITE FASCICULATION, NEURITES, ETC. STANDARD; PRT; 1088 (shown here) and IN PRESUMPTIVE NEURAL DOMAINS. **EMBRYO** THE MAJOR DURING

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Alignment Scores:
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pfam; pF00047; ig; 5.
SMART; SM00060; FN3; 2.
SMART; SM00408; IGC2; 5.
Cell adhesion; Glycoprotein; T.
Immunoglobulin domain; Alterna
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InterPro; IPR003006; Ig_MHC
InterPro; IPR003598; Ig_c2.
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                                 {\tt IleSerPheAsnGluAspGlnSerGluMetThrIleHisHisValGluLysAspAspGluInterval} \\
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Eukaryota; Metazoa; (Mammalia; Eutheria; (NCBI_TaxID=9823;
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Q9GL76;
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence up
15-JUN-2002 (Rel. 41, Last annotation
15-JUN-2002 (Rel. 41, Last annotation
                              MEDLINE=21176378; PubMed=11277703; Milne R.S.B., Connolly S.A., Krummv Cohen G.H.;
                                                                                                                                                 Poliovirus receptor related mediator C) (HveC) (Nectin PVRL1 OR PRR1 OR HVEC.
is a functional alphaherpesvirus Virology 281:315-328(2001).
                      "Porcine
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                       a member of the
                                                                                                            Chordata; Craniata; Vertebrata; Cetartiodactyla; Suina; Suidae;
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Minimum DB:
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MODEL-frame+_n2p.model -DEV=xlp
-Q-cgp2_1/USPTO_Spool_US09778187/runat_20112002_073631_26345/app_query.fasta_1.2318
-DB-Issucd_Patents_AA -OFMT-fastan -SUFFIX=rai -MINMATCH=0.1 -LOOPCL=0
-LOOPEXY=0 -UNITS-blts -STARY=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
-LIST=45 -DCCALIGN=200 -THR_SCORE=pot -THR_MAX=100 -THR_NIN=0 -ALIGN=15
-MODE=LOCAL -OUTFMT-pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
-USER=US09778187_eCm_1_1_28_grunat_20112002_073631_26345 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MAMP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
-VGAPOP=10 -VGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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/cgn2_6/ptodata/1/18a/5B_COMB.pep:*
/cgn2_6/ptodata/1/18a/6A_COMB.pep:*
/cgn2_6/ptodata/1/18a/6B_COMB.pep:*
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US-08-429-742-2
US-08-429-742-4
US-08-217-299-1
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US-08-659-984A-1
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### ALIGNMENTS

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Sequence 1, Application US/08659984A
Patent No. 5942400
GENERAL INFORMATION:
APPLICANT: Anderson, John P.
APPLICANT: Sinha, Sukanto
APPLICANT: Jacobson-Croak, Kirsten L.
TITLE OF INVENTION: Assays for Detecting Beta-Secretase
TITLE OF INVENTION: Inhibition
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Ctr., 8th Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 9411-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US.08/485.152
FILING DATE: 07-JUN-1996
CLASSIFICATION NUMBER: US.08/485.152
FILING DATE: 07-JUN-1996
APPLICATION NUMBER: US.08/485.152
FILING DATE: 07-JUN-1996
APPLICATION NUMBER: US.08/485.152
FILING DATE: 07-JUN-1996
REGERENCE/DOCKET NUMBER: 15270-002810US
TELECOMMUNICATION INFORMATION:
REGISTRATION NUMBER: 15270-002810US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: sir
TOPOLOGY: linear
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CTGTTCATCAATAACCTAAACAAAACAGATAATGGTACATACCGCTGTGAAGCTTCAAAC
                         TrpThrLysAspGlyGlyGluLeuProAspProAspArgMetValValSerGlyArgGlu
                                                           TGGGTGAGAGTCGATGAAATG-----CCTCAACACGCCGTACTGTCTGGGCCCAAC
                                                                                       GlyGlnProLeuIleLeuThrCysGluSerLysGlyLysProLeuProGluProValLeu
                                                                                                                                                                                 CAGTATAAGCCTCAAGTGCACATTCAGATGACTTATCCTCTACAAGGCTTAACCCGGGAA
                                                                                                                                                                                                                                              CAGGTGGAGCACCCTGCGGTCACTGGAAACCTGCAG----ACCCAGCGGTATCTAGAAGTA
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                                                                                                                      GGGGACGCGCTTGAGTTAACATGTGAAGCCATCGGGAAGCCCCCAGCCTGTGATGGTAACT
                                                                                                                                                    HisTyrThrProSerValLysIle-----IleProSerThrProPheProGlnGlu
                                                                                                                                                                                                                 ArgValAspHisGluSerLeuAsnAlaThrProGlnValAlaMetGlnValLeuGluIle
                                                                                                                                                                                                                                                                                                         GTGACCAGTCAGCTGATGCTGAAGGTGCACAAGGAGGACGATGGGGTCCCAGTGATCTGC
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Query Match:
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                         Percent Similarity:
Best Local Similarity:
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US-08-660-531-1
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US-08-660-531-1
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                                                        Score:
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                                                                                                                                                                                                           TELEFAX: 415-326-2422 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 0 FILING DATE: 07-JUN-1995 ATTORNEY/AGENT INFORMATION:
                                                                   No.:
                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 421 amino acids
                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: 415-326-2400
                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
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ADDRESSEE: Townsend and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Chrysler,
APPLICANT: Sinha, Su
APPLICANT: Keim, Pam
APPLICANT: Anderson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
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OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                             NAME: Heslin, James M. REGISTRATION NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: San Francisco
STATE: California
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PURNTION: Beta-Secretase
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Result
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-MODEL-frame-n2p.model -DEV-xlp
-O-/cgn2_1/USPTO_spool/US09778187/runat_20112002_073630_26319/app_query.fasta_1.2318
-O-/cgn2_1/USPTO_spool/US09778187/runat_20112002_073630_26319/app_query.fasta_1.2318
-DB-SPTREMBL_21 -OFMT-fastan -SUFFIX-rspt -MINMATCH-0.1 -LOOPCL-0 -LOOPEXT-0
-UNITS-bits -START-1 -END--1 -MATRIX-blosum62 -TRANS-human40.cd1 -LIST-45
-DOCALIGN-200 -THR_SCORE-pct -THR_MAX=100 -THR_MIN-0 -ALIGN-15 -MODE-LOCAL
-OUTFMT-pto -NORM-ext -HEAPSIZE-500 -MINILEN-0 -MAXLEN-2000000000
-USER-US09778187_eCGN_1_1_204_@runat_20112002_073630_26319 -NCPU-6 -ICPU-3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES-0 -WAIT -LONGLOG -DEV_TIMEOUT-120
-MARN_TIMEOUT-30 -THREADS-1 -XGAPD-10 -XGAPEXT-0.5 -FGAPEXT-7
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### ALIGNMENTS

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RESULT 1
QSBY67
QSBY67
ID QSBY67
AC QSBY67
PT 01-JUN-2001 (TrEMBLrel. 17, Created)
DT 01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
DT 01-JUN-2001 (TrEMBLrel. 19, Last annotation update)
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
DE Nectin-like protein 2.
GN NECL2.
GN NECL2.
GN NECL2.
OC Enkaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OC NCBL_TAXID=9606;
RN [1]
RN [1]
RN SEQUENCE FROM N.A.
RP SEQUENCE FROM N.A.
RP SEQUENCE FROM N.A.
RP SEQUENCE FROM N.A.
Thou Y., Du G., Chen J., Yuan J., Qiang B.;
RN [1]
RN [1]
RN [1]
RN [1]
RN [1]
RN [1]
RN [2]
Submitted (MaR-1999) to the EMBL/GenBank/DDBJ databases.
DR InterPro; IPR003598; Ig_
DR InterPro; IPR0035
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Alignment Scores:
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RESULT
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Q8R4L1;
Q1-JUN-2002 (TrEMBLrel. 21, Created)
Q1-JUN-2002 (TrEMBLrel. 21, Last sequence update)
Q1-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Tumor suprressor in lung cancer 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.

STRAIN-129/SVJ;

FUKAMI T., MARLYDAMA T., MURAKAMI Y.;

FUKAMI T., MARLYDAMA T., MURAKAMI Y.;

"Identification of murine orthologue of the TSLC1 gene."

SUBMILTED (CCT-2001) to the EMBL/GenBank/DDBJ databases.

EMBL; AF434663; AAL86736.1; -.

SEQUENCE 445 AA; 48664 MW; C5D5A070DAF70E55 CRC64;
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10090;
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"Blological function of RA175,
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Submitted (JUN-2001) to the EMB
EMBL; AB064265; BAB83501.2;
SEQUENCE 456 AA; 49787 MW;
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Eukaryota; Metazoa; Chordata;
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Arakawa T., Hara A., Pukunishi Y., Konno H., Adachi J., Fukuda S.,
RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
RA Fleischmann W., Gaasterland T., Glssi C., King B., Kochiwa H.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Jee N.H.,
Lyons P., Marchionni L., Mashima J., Mazzarelli J., Mombaerts P.,
RA Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,
RA Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,
RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,
RA Wynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
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7BL/6J; TISSUE=EMBRYONIC |
1085660; PubMed=11217851;
Shinagawa A., Shibata K.
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95.68%
95.81%
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Rodentia;
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Q9D6E7;
Q1-JUN-2001;
01-JUN-2001;
01-DEC-2001;
                                    Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10090;
[1]
SEQUENCE FROM N.A. STRAIN-C57BL/6J; TISSUE-HIPPOCAMPUS; MEDLINE-2108560; PubMed-11217851; Kawai J., Shinagawa A., Shibata K.,
                                                                               01-JUN-2001 (TrEMBLrel.
01-DEC-2001 (TrEMBLrel.
2900073G06Rik protein.
IGSF4 OR 2900073G06RIK.
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RA AIZBWA K., IZBWA M., NIShi K., Kiyosawa H., Kondo S., Yamanaka I., RA AIZBWA K., IZBWA M., NIShi K., Kiyosawa H., Kondo S., Yamanaka I., RA Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R., RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T., RA Kuchi P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J., RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T., RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G., RA Slakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G., RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F., RA Blake J., Bult C., Fletcher C., Fujita M., Gariboldi M., RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H., RA Lyons P., Marchionni L., Mashima J., Mazzarelli J., Wombaerts P., RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F., RA Sasaki H., Sorok K., Wang K.H., Weltz C., Whittaker C., wilming L., RA Hygashizaki Y., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S., RT Paritana.
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Best Local Similarity:
Query Match:
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InterPro; IPR003598; Ig_C2.
InterPro; IPR003500; Ig_like.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003006; Ig_MHC.
INTERPRO; IG; 3.
SMART; SM00409; IG; 3.
SMART; SM00408; IGC2; 3.
SMART; SM00410; IG_Like; 1.
Immunoglobulin domain.
SEQUENCE 336 AA; 37157 MW;
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EMBL; AKO13775; BAB28988.1; -.
MGD; MGI:1889272; Igsf4.
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99.32%
84.90%
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A Zhou Y., Du G., Chen J., Yuan J., Qlang B.;

A Zhou Y., Du G., Chen J., Yuan J., Qlang B.;

T "Cloning of a novel cDNA encoding a member of inm

Submitted (APR-1998) to the EMBL/GenBank/DDBJ dat

R EMBL; AF061260; AAC67243.1; -.

MG]: MG]: 1889272; Igsf4.

R InterPro; IPR003598; Ig_C2.

R InterPro; IPR003500; Ig_like.

InterPro; IPR003600; Ig_HKC.

R InterPro; IPR003585; Neurexin-like.

R InterPro; IPR003585; Neurexin-like.

R Ffam; PF00047; 19; 2.

R SMART; SM00408; IGC2; 1.

R SMART; SM00408; IGC2; 1.

R SMART; SM00408; IGC2; 1.

Immunoglobulin domain.

SEQUENCE 295 AA; 32509 MW; 9DE9D86F6FF6F488 C
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Q9Z2H8;
01-MAY-1999
01-MAY-1999
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Nectin-like protein 2.
IGSF4 OR NECL2.
                                                                                                                                                                                                                                                                                                       Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10090;
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InterPro; IPR003600; Ig_like.
InterPro; IPR003006; Ig_like.
InterPro; IPR003085; Neurexin-like.
InterPro; IPR003585; Neurexin-like.
InterPro; IPR003585; Neurexin-like.
InterPro; IPR003585; Igc2; 1.
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Q9QYL4;
Q1-MAY-2000 (Treme
01-MAY-2000 (Treme
01-JUN-2002 (Treme
01-JUN-2002 (Treme
Adhesion protein A
                                                                                                        "RA175, a novel neuron spec:
Submitted (DEC-1998) to the
EMBL; AB021966; BAA87916.1;
MGD; MGI:1889272; Igsf4.
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Eukaryota; Me
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SEQUENCE 306 A
Eukaryota;
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AsnGlyThrTyrProCysGluAlaSerAsnIleValGlyLysAlaHisSerAspTyrIle
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            ion protein OR RA175A.
Metazoa;
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cotein RA175A.
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Vertebrata;
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SMART; SM00294; 4.1m; 1.
SMART; SM00408; IGc2; 1.
SMART; SM00410; IG_like; 1.
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InterPro;
InterPro;
InterPro;
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NCBI_TaxID=10090;
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                 CCTCAACACGCGGTACTGTCTGGGCCCAACCTGTTCATCAATAACCTAAACAAAACAGAT
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IPR003600; Ig_like.
IPR003006; Ig_MHC.
IPR003585; Neurexin-like.
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Fujita E.,
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"RA175, a
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SMART; SM00294; 4.1m; 1.
SMART; SM00408; IGC2; 1.
SMART; SM00410; IG_11ke;
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InterPro; IPR003598; Ig_C2.
InterPro; IPR003600; Ig_HLC.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003585; Neurexin-like.
Pfam; PF00047; 19; 2.
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Submitted (DEC-1998) to the
EMBL; AB021955; BAA87915.1;
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SEQUENCE 289
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Mammalia; Eutheria;
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Pfam; PF00047; 19; 2.
SMART; SM00294; 4.1m; 1.
SMART; SM00408; IGC2; 1.
SMART; SM00410; IG_like; 1.
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MGD; MGI:1889272; Igsf4
InterPro: Tppc?-
                                                                                                                                                        Immunoglobulin
SEQUENCE 278
                                                                                                                                                                                                                                                                                   Soyama
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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01-MAY-2000 (TrEMBLrel. 13,
01-MAY-2000 (TrEMBLrel. 13,
01-JUN-2002 (TrEMBLrel. 21,
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InterPro; IPR003600; Ig_like.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003585; Neurexin-like.
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01-JUN-2002
01-JUN-2002
01-JUN-2002
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"Identification of a murine ortholog of the TSLC1-like of Submitted (OCT-2001) to the EMBL/GenBank/DDBJ databases EMBL; AY059394; AAL29692.1;
                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A. STRAIN=129/SVJ;
                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=10090;
                                                                                                                                                                                                                       SEQUENCE
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Sciurognathi; Muridae;
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Primates;

Catarrhini; Hominidae;

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RESULT
Q9Y4A4
ID Q9
AC Q9
DT 01
DT 01
DT 01
DT 01
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OS HO
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                     O9Y4A4 PRELIMINARY;

O9Y4A4;

O1-NOV-1999 (TrEMBLrel. 12, C

O1-NOV-1999 (TrEMBLrel. 12, L

O1-JUN-2002 (TrEMBLrel. 21, L

F2216_1 (Fragment).
               Homo sapiens
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   Metazoa;
               (Human)
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Burkhart-Schultz K.J., Gordon L., Kyle A., Ramirez M., Stilwagen S.,
Phan H., Velasco N., Do L., Regala W., Terry A., Garnes J.,
Danganan L., Poundstone P., Christensen M., Georgescu A., Avila J.,
Liu S., Attix C., Andreise T., Trankheim M., Amico-Keller G.,
Coeffeld J., Duarte S., Lucas S., Bruce R., Thomas P., Quan G.,
Kronmiller B., Arellano A., Montgomery M., Ow D., Nolan M., Trong S.
Kronmiler B., Arellano A., Montgomery M., Ow D., Nolan M., Trong S.
Krobayashi A., Olsen A.S., Carrano A.V.;
Sequence analysis of a 2.5 Mb region in 19q13.2 containing a
clustered (EMAPSG gene family "
Submitted (AUG-1998) to the EMBL/GenBank/DDBJ databases.
EMBL, ACO05525, AAC32740.1; -.
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SEQUENCE
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InterPro; IPRO
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                                                                                                                                                                           CCACGTAATCTGATGATCGATATCCAGAAAGACACTGCGGTGGAAGGTGAAGGAGATTGAA
                                                                                                                                                                                                                    CysGlnLeuTyrThrGluAspThrHisHisGlnIleAlaThrLeuThrValLeuValAla 101
                                                                                                                                                                                                                                                                                                                                                                                                                           TGCCAAGTCAATAAGAGTGACGACTCTGTGATTCAGCTACTGAATCCCAACAGGCAGACC
                                       Lys \verb|GluLeuLysGlyValSerSerSerGlnGluAsnGlyLysValTrpSerValAlaSer|
                                                                     ACAGAGCTAAAAGGCAAATCGGAGGTGGAAGAGTGGTCAGACATGTACACTGTGACCAGT
                                                                                                  LeuSerCysLeuValProArgSerArgProAlaAlaThrLeuArgTrpTyrArgAspArg
                                                                                                                                                            ProGluAsnProValValGluVal---ArgGluGlnAlaValGluGlyGlyGluValGlu
                                                                                                                                                                                                                                                                           {\tt ProArgArgValArgIleArgLeuSerAspAlaArgLeuGluAspGluGlyGlyTyrPhe}
                                                                                                                                                                                                                                                                                               SM00408;
SM00410;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IPR003598; Ig_c2.
IPR003600; Ig_like.
IPR003006; Ig_MHC.
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IG_11ke;
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58.39%
39.13%
33.81%
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Matches:
Conservative:
Mismatches:
Indels:
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                                 SMART;
                                                                                                                                        EMBL;
                                                                                                                                                Fukami T., Maruyama T., Murakami Y.;
"Identification of a murine ortholog of the T.
"Submitted (OCT-2001) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                   "Cloning membrane
                                                                                                                                                                                                                                                                                                                                            099N28 PRELIMINARY; PRT; 396 AA. 099N28; 01-JUN-2001 (TrEMBLrel. 17, Created) 01-JUN-2001 (TrEMBLrel. 17, Last sequence update) 01-JUN-2002 (TrEMBLrel. 21, Last annotation update) Nectin-like protein 1 (Membrane glycoprotein).
                                                                                                                                                                                                                                                                                                  Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                     SMART; SM00408; IGC2; 3. SMART; SM00410; IG_like;
                                                                                                                  MGD; MGI:2137858;
                                                                                                       InterPro; IPR003599;
                                                                                                                                                                                                                                 Zhou Y., Huang X., Yuan
"Cloning and expression
membrane protein.";
                                                        Pfam; PF00047; ig;
                                                                     InterPro;
                                                                                InterPro;
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                                                                                                                                                                                     STRAIN=129/SVJ;
                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                         NCBI_TaxID=10090;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    {\tt Thr LeuVal LeuThr CysAlaVal Thr Gly Asn Pro Arg Pro Asn Gln Ile Arg Trp Asn}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATCCTT
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                                                                                                                          AF195662; AAG35584.1;
AY059393; AAL29691.1;
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IPR003600;
                                                                    IPR003006;
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                                                      {\tt AlaThrSerAsnMetGlySerTyrThrAlaTyrPheThrLeuAsnValAsnAspProSer}
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InterPro: IPR003600; Ig_llke.
InterPro: IPR003006; Ig_MHC.
InterPro: IPR003585; Neurexin-like.
Pfam: PF00047; 1g; 3
SMART; SM00294; 4.1m; 1.
SMART; SM00294; 4.1m; 1.
SMART; SM00400; IGc2; 1.
SMART; SM00410; IG_llke; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Zhou Y., Du G., Yuan J., Qiang B.; Submitted (APR-2001) to the EMBL/GenBank/DDBJ EMBL; AL035403; CAB56227.1; -. EMBL; AF062733; AAD17540.2; -.
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
BK134P22.1 (Novel protein similar to mouse IMMUNOSUPERFAMILY protein
BL2) (NECTIN-11ke protein 1).
BK134P22.1 OR NECTL.
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Submitted (AUG-1999)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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SerSerGlySerLysProAlaAlaArgLeuThrTrpArgLysGlyAspGlnGluLeuHis
                                              GCTATGGCCAGCCAGCCACGACTATCAGGTGGTTCAAAGGGAACACAGAGCTAAAA
                                                                                                                                            ATGATCGATATCCAGAAAGACACTGCGGTGGAAGGTGAGGAGATTGAAGTCAACTGCACT
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                              RA Kawai J. Shinagawa A. Shibata K. Yoshino M., Itoh M., Ishii Y., Ra Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S., RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S., RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I., RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I., RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I., RA Raito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R., RA Raito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R., RA Raito T., Fuzuna M., Anno H., Baitaloy S., Casavant T., RA Schriml L.M., Stabbli F., Suzuki R., Tomita M., Wagner L., Washio T., RA Schriml L.M., Stabbli F., Suzuki R., Tomita M., Garibodi M., RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G., RA Bake J., Boffeli D., Bojunga N., Carninci P., de Bonaldo M.F., RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Garibodi M., RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H., RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H., RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H., RA Gustincich S., Hill D., Hofmann M., Bodriguez I., Sakamoto N., Schoenbach C., Seya T., Shibata Y., Storch K.-F., RA Suzuki H., Toyo-oka K., Wang K.H., Weitz C., Whittaker C., Wilming L., Washaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S., Landar M., Kohtsuki S., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
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Q9D006;
01-JUN-2001 (TrEMBLrel. 1
01-JUN-2001 (TrEMBLrel. 1
01-DEC-2001 (TrEMBLrel. 1
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STRAIN-C57BL/6J; TISSUE-EMBRYO;
MEDLINE-21085660; PubMed-112.7851;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus (Mouse).
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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EMBL; AK011949; BAB27933.1; -.
MGD; MGI:1914402; 26130301B19Rik.
InterPro; IPR003599; Ig.
InterPro; IPR003599; Ig__1ike.
InterPro; IPR003006; Ig__1ike.
InterPro; IPR0030006; Ig__MHC.
Pfam; PF00047; ig; 2.
SMART; SM00410; IG; 1.
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SMART; SM00410; IG; 1.
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294 AspAlaAsnProProProPheLysSerValTrpSerArgLeuAspGlyGlnTrpProAsp
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                ATCGGGAAGCCCCAGCCTGTGATGGTAACTTGGGTGAGAGTCGATGATGAAATGCCTCAA
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                                                             TyrAspGlyAsnTrpPheValGlyArgLysGly-----ValAsnLeuLysCysAsnAla
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Search completed: November 20, 2002, 07:55:59 Job time: 76.25 secs

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GenCore version 5.1.3 Copyright (c) 1993 - 2002 Compugen Ltd.
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Command line parameters:

MODEL-frame+_n2p.model -DEV-xlp
-MODEL-frame+_n2p.model -DEV-xlp
-O_/Ggn2_1/USPTO_Spool_US09778187/runat_20112002_073629_26303/app_guery.fasta_1.2318
-OB-A_Geneseq_101002 -OFMT-fastan -SUFFIX-rag -MINMATCH-0.1 -LOOPCL-0
-LOOPEXY=0 -UNITS-blts -STARY=1 -END=-1 -MATRIX-blosum62 -TRANS-human40.cdi
-LIST-45 -DOCALIGH-200 -THR_SCORE-pot -THR_MAX-100 -THR_NIN-0 -ALICNA-15
-NODE-LOCAL -OUTFMT-pto -NORM-ext -HEAPSIZE-500 -MINLEN-0 -MAXLEN-200000000
-USER-US09779187_@CGN_1_1_71 @runat_20112002_073629_26303 -NCPU-6 -ICPU-3
-NO_MARP -LARGEQUERY -NEC_SCORES-0 -WAIT -LOULOG -DEV_THREOUT-120
-WARN_TIMEOUT-30 -THREADS-1 -XGAPOP-10 -XGAPEXT-0.5 -FGAPOP-6 -FGAPEXT-7
-YGAPOP-10 -YGAPEXT-0.5 -DELOP-6 -DELEXT-7
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

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## ALIGNMENTS

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AAM23691;

12-OCT-2001 (first entry)

Human EST encoded protein SEQ ID NO: 1216.

Human; sheep; plg; cow; fruit fly; yeast; hamster; macaque; tomato; monkey; dog; sea urchin; expressed sequence tag; ES? diagnostics; forensic test; gene mapping; genetic disorder; biodiversity; gene therapy; nutrition. norse;

Homo sapiens.

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N-PSDB;
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Hillan KJ,
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16-DEC-1998;
22-DEC-1998;
                                                                                                                                                                                                                       New human nucleic acids encoding secreted and transmembrane polypeptides which are designated as PRO polypeptides are described The membrane-bound proteins have various industrial applications, including as pharmaceutical and diagnostic agents. The membrane-bound proteins can also be employed for screening of potential peptide or small molecule inhibitors of the relevant receptor/ligand interaction. Anti-PRO antibodies are useful for the affinity purification of PRO from recombinant cell culture or natural sources.
                                                                                                                                                                                                                                                                                                           Claim 12;
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ci PJ, Grimaldi CJ, Gurney
L, Roy MA, Tumas D, Wood W
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                                   t; rabbit; tumour necrosis factor alpha; TNF-alpha;
cell; cell proliferation; cell differentiation; colon;
ast; prostate; rectum; cervix; liver; genetic disorder.
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25-APR-2000

27-AUG-2000

28-AUG-2000

28-AUG-2000

29-AUG-2000

20-DEC-2000

20-DEC-2000
Sequences AAU29024-AAU29328 represent PRO polypeptides of the invention. The PRO polypeptides and their associated nucleic acids can be used to detect the presence of a tumour in a mammal by comparing the level of expression of a PRO polypeptide in a test sample of cells from the animal and a control sample of normal cells, whereby a higher level of expression in the test sample indicates the presence of a tumour in the mammal include dogs, cats, cattle, horses, sheep, pigs, goats and rabbits but are preferably human. The polypeptides can be used to stimulate tumour necrosis factor (TR) alpha release from human blood, when contacted with it. A specific polypeptide can be used to stimulate the proliferation or differentiation of chondrocyte cells. The PRO proteins can be used to determine the presence of tumours and also
                                                                                                                                                                        Novel nucleic acids encoding PRO polypeptides, used presence of tumours, such as prostate and breast tun to screen for modulators of the compounds -
                                                                                                                                                     Claim
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                                            ACTTGGGTGAGAGTCGATGATGAATGCCTCAACACGCCGTACTGTCTGGGCCCAACCTG
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The present invention relates to 12 secreted human proteins and the CC nucleotide sequences encoding them. The polynucleotide sequences given in AAA80606-A80623 encode the 12 secreted protein sequences given in CC AAB2576-B2593. The human secreted proteins have various activities CC dependent on the tissues in which they are expressed. Examples of the cativities of the proteins include: immunosuppressant; CC antirinflammatory; antiarthritic; antirheumatic, dermatological; CC antivital; antiarteriosclerotic; anticancer; vulnerary; CC antivital; antibacterial; and antifungal activity. The proteins, CC polypeptides, agonists and antagonists may be used to treat prevent CC and/or diagnose various disease, disorders and conditions examples of CC which include: immune disorders e.g. Addison's disease, rheumatoid CC arthritis, dermatitis, and multiple sclerosis; inflammatory disorders ce.g. inflammatory bowel disease, Crohn's disease and nephritis; CC hyperproliferative disorders such as paraproteinaemias and purpura; CC cardiovascular disorders e.g. coronary arteriosclerosis and myocarditis; cancer e.g. melanoma and lymphoma. The proteins and polynucleotide constructions and all molynucleotide sequences may also be used in wound healing and the treatment of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Isolated nucleic acid molecules encoding human secreted proteins a used for the prevention, amelioration and treatment of autoimmune, inflammatory, hyperproliferative and cardiovascular disorders, can wounds, and infectious diseases.
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                                                                                                                                                                                                                                                                                                                                       disorders such as rheumatoid arthritis, asthma and atherosclerosis, bacterial and parasitic infections and neuronal disorders such as akathesia, Alzheimer's disease, multiple sclerosis and epilepsy. Polynucleotides encoding HSCRPs may be used as hybridisation probes diagnose these conditions. Anti-HCSRP antibodies may be used as antagonists, as a targeting or delivery mechanism for bringing pharmaceutical agents into contact with cells or tissues expressing HCSRP and for diagnosis of HCSRP-related disorders. HCSRP and its
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07-DEC-1998;
08-MAR-1999;
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                                                                                                                                                                                                                                                                                      Sequence
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organ

transplant rejection;

immune system disorder; autoimmune; viral infection; inf

bone

marrow;

modulator; immune response

Best Local Similarity:
Query Match:
DB:

Percent Similarity:

7.82e-151 1761.00 100.00% 100.00% 100.00% 97.78%

Length:
Matches:
Conservative:
Mismatches:
Indels:

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Alignment Scores:

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infectious disease

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The present amino acid sequence is the human lymphoid derived dendritic CC cells adhesion molecule, LDCAM. It is found on lymphoid derived dendritic CC cells and displays homology to adhesion molecules. B7-1 and cytoplasmic CC region of B7-1. Human LDCAM is expressed in breast, retina, foetal C1 liver, spleen and heart, lung, muscle, placenta, thyroid and lung CC carcinoma. LDCAM polypeptides interacts with T cell surface molecules CC to alter signalling and inhibits T cell proliferation, bind to CC themselves and B7L-1, an LDCAM binding protein and increases natural CC killer (NK) cell populations. It may be used to measure the biological CC activity and as quality control reagents of LDCAM binding proteins. CC LDCAM may be used for treating disorders associated with malfunctioning CC immune system, inflammation, autolimnune disorders, viral infected CC cells, infectious diseases and for killing tumour cells. They are also CC useful for prevention or reducing the effect of organ and bone marrow CC useful for prevention and for modulating T cell immune responses. LDCAM CC polypeptides may also be used as carriers for delivering agents attached CC cells or cells bearing B7L-1.
Sequence
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Human; hepatocellular carcinoma; tumour suppressor liver; lung; pancreatic cancer; cell proliferative
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                                                     cancer 1 (TSLC1)
                                                     polypeptide
   lung canc
disorder;
                 cancer 1;
   er 1; TSLC1; cytostatic;
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Detecting cell proliferative disorder associated with tumor suppressor lung cancer (TSLC) 1 in subject, comprises contacting proliferating cell of subject with reagent detecting TSLC1 and detecting modification in TSLC1 level -

English

The invention relates to a method for detecting cell proliferative disorder associated with tumour suppressor lung cancer 1 (TSLC1) in a subject. The method comprising contacting a cell component of a proliferating cell with a reagent that detects level of the cell component in the proliferating cell and determining modification in the level of the cell component in proliferating cell as compared with a healthy cell, where modification indicates disorder associated with TSLC1. The method is useful for detecting a cell proliferative disorder (e.g. liver, lung or pancreatic cancer) associated with tumour suppressor lung cancer 1 (TSLC1) in a subject. The invention is useful in gene therapy and for treating a cell proliferative disorder such as lung cancer (human non-small cell lung cancer), liver cancer (hepatocellular cancer (human non-small cell lung cancer), liver cancer (hepatocellular cancer of modulates) are cancer a reagent which modulates (preferably, increases) TSLC1 production, where a reagent which modulates (preferably, increases) TSLC1

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Length:
Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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- GTTGCGACCATCAGTTGCCAAGTCAATAAGAGTGACGACTCTGTGAGTTCAGCTACTGAAT
- CCCAACAGGCAGACCATTTATTTCAGGGACTTCAGGCCTTTGAAGGACAGCAGGTTTCAG

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       Lymphoid derived dendritic cell adhesion molecule; LDCAM; mouse; B7-1; B7-L1; T cell proliferation; natural killer cell; NK; tunnour cell; biological activity; quality control reagent; treatment; inflammation; immune system disorder; autoimmune; viral infection; infectious disease;
                                                       Mouse
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Percent Similarity: Best Local Similarity:

4.99e-149 1741.00 99.11% 98.81% 96.67%

Conservative: Mismatches: Indels:

Length: Matches:

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The present amino acid sequence is the mouse lymphoid derived dendrtitic CC cells and displays homology to adhesion molecules, B7-1 and cytoplasmic CC cells and displays homology to adhesion molecules, B7-1 and cytoplasmic CC region of B7-L1. Mouse LDCAM is found on whole embryo, testes, triple CC negative cells murine splenic and lymph node CD8+, S49.1 and dendritic CC cells. LDCAM polypeptides interacts with T cell surface molecules CC to alter signalling and inhibits T cell proliferation, bind to CC themselves and B7L-1, an LDCAM binding protein and increases natural CC killer (NK) cell populations. It may be used to measure the biological activity and as quality control reagents of LDCAM binding proteins. CC LDCAM may be used for treating disorders associated with malfunctioning CC cells, infectious diseases and for killing tumour cells. They are also useful for prevention or reducing the effect of organ and bone marrow cc useful for prevention and for modulating T cell immune responses. LDCAM college may also be used as carriers for delivering agents attached
                          polypeptides may also be used
to T cells or cells bearing BJ
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The present invention relates to 12 secreted human proteins and the concleotide sequences encoding them. The polynucleotide sequences given in AAA80606-A80623 encode the 12 secreted protein sequences given in CC AAB2576-B2593. The human secreted proteins have various activities of the tissues in which they are expressed. Examples of the cativities of the proteins include: immunosuppressant; antiplicative; antiarthrilic; antirheumatic, dermatological; antiproliferative; antiarteriosclerotic; anticancer; vulnerary; cc antivital; antibacterial; and antifungal activity. The proteins; cc antivital; antibacterial; and antifungal activity. The proteins of the polypeptides, agonists and antigonists may be used to treat prevent and/or diagnose various disease, disorders and conditions examples of which include: immune disorders e.g. Addison's disease, rheumatoid arthritis, dermatitis, and multiple sclerosis; inflammatory disorders e.g. Addison's disease and nephritis; thyperproliferative disorders such as paraproteinaemates and purpura; cardiovascular disorders e.g. coronary arteriosclerosis and myocarditis; cancer e.g. melanoma and lymphoma. The proteins and polynucleotide approaches may also be used in wound healing and the treatment of a first and polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Secreted protein; immunosuppressant; anti-inflammatory; antiarthritic; antirheumatic, dermatological; antiproliferative; antiarteriosclerotic; anticancer; vulnerary; antiviral; antibacterial; antifungal; immune disorder; Addison's disease; rheumatoid arthritis; dermatitis; multiple sclerosis; inflammatory disorder; inflammatory bowel disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Crohn's disease; nephritis; hyperproliferative disorder; cancer; cardiovascular disorder; coronary arteriosclerosis; myocarditis; cancer;
                                   infectious diseases. The human secreted protein gene #11 and protein sequences are represented in sequences AAA80616 and AAB25586. Sequen AAA80677-A80682 represent genes related to the secreted protein gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolated nucleic acid molecules encoding human secreted proteins are used for the prevention, amelioration and treatment of autoimmune, inflammatory, hyperproliferative and cardiovascular disorders, cancel wounds, and infectious diseases.
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N-PSDB; AAA80616.
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                                                                                                                                                                                       (e.g. polymerase chain reactions (PCR)) to detect and quantitate the presence of similar nucleic acid sequences in samples. They may also be used to study the expression and function of secretory proteins/membrane polypeptides and their role in metabolism. The polypeptides may be used as antigens in the production of antibodies against them and in assays to identify modulators (agonists and antagonists) of expression and activity. The antibodies and antagonists may also be used as therapeutic agents to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of the polypeptides in samples (e.g. by enzyme linked immunosorbant assay (ELISA). Examples of diseases which may be treated include rheumatoid arthritis and diabetes.
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11-JAN-2000; 2
02-MAY-2000; 2
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                                                                                  GTGGGGAAAGCTCACTCGGATTATATGCTGTATGTATACGATCCCCCCACAACTATCCCT
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                                                                                                   ATCACAGATTCCCGAGCAGGTGAAGAAGGCTCGATCAGGGCAGTGGATCAT 1008
                                                                                                                    ProAlaValHisGlyLeuThrGlnLeuProAsnSerAlaGluGluLeuAspSerGluAsp
                                                                                                                                     ValGlyLysAlaH1sSerAspTyrMetLeuTyrValTyrAspThrThrAlaThrThrGlu
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Best Local Similarity:
                                               US-09-778-187B-1_COPY_130_1137
                                                                                   Query Match:
                                                                                                                        Score:
                                                                                                                                                                                                   The present invention describes new human secreted proteins which were isolated from adult placenta, adult retina, foetal brain, foetal kidney, adult blood, adult blood, adult train, adult tymph node cDNA libraries. The human secreted proteins, and the polynucleotides encoding them, are predicted to have biological activities which would make them suitable for treating, preventing or ameliorating medical conditions in humans and animals. Suggested activities include nutritional activity, cytokine and cell proliferation/differentiation activity, haematopolesis regulating (e.g. as vaccines) or suppressing activity, haematopolesis regulating activity, tissue growth activity, activity, haematory activity, cactivity, receptor/ligand activity, activity, and thrombolytic activity. The polynucleotides are also stated to be useful for gene tharapy. AAZ33316 to AAZ33373 encode human secreted proteins, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-JUL-1998;
31-JUL-1998;
10-AUG-1998;
11-AUG-1998;
06-MAY-1999;
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differentiation; immune stimulating; vaccine; suppression;
haematopoiesis regulation; tissue growth; activin; inhibin;
chemotactic; chemokinetic; haemostatic; thrombolytic; receptor;
ligand; anti-inflammatory; cadherin; tumour invasion suppressor;
tumour inhibition; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                               Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                     New polynucleotides encoding secreted human proteins, adult placenta, adult retina, fetal brain, fetal -
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DB; AAZ33346.
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Conservative:
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RESULT 1:
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ID AAB:
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                      21-NOV-2000
                                             AAB25593;
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                                                                  standard;
                      (first entry)
by human secreted protein gene
                                                                  Protein;
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 #11 clone HOUDJ81.
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Crohn's disease; nephritis; hyperproliferative disorder; cardiovascular disorder; coronary arteriosclerosis; myoc
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                                                                                                                                                                                                                                                                                                                    28-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                    WO200029435-A1
                                                                                                                                                                                                                                                                                                                                                                                                    melanoma; lymphoma; wound healing; human
                                                                                                                                                                                                                                                                                                                                                                                                                            multiple sclerosis; inflammatory disorder; inflammatory bowel
                                                                                                                                                                                                                                                                                                                                    27-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                    99WO-US25031
                                                                                                                                                                                                                                                                                                                                                                                                            myocarditis; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                             disease;
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Isolated nucleic acid molecules encoding human secreted proteins a used for the prevention, amelioration and treatment of autoimmune, inflammatory, hyperproliferative and cardiovascular disorders, can wounds, and infectious diseases. WPI; 2000-387742/33 Greene Ni J, Ruben SM, Olsen HS, Young PE, Kenny JJ, Moore Wei are ĸ

(HUMA-) HUMAN GENOME SCI INC

Claim 1; Page 685-686; 803pp; English.

anti-inflammatory; antiarthritic; antirheumatic, dermatological; antiproliferative; antiarteriosclerotic; anticancer; vulnerary; antiviral; antibacterial; and antifungal activity. The proteins, polypeptides, agonists and antagonists may be used to treat prevent and/or diagnose various disease, disorders and conditions examples of which include: immune disorders e.g. Addison's disease, rheumatoid arthritis, dermatitis, and multiple sclerosis; inflammatory disorders e.g. inflammatory bowel disease, Crohn's disease and nephritis; e.g. inflammatory bowel disease, Crohn's disease and nephritis; hyperproliferative disorders such as paraproteinaemias and purpura; cardiovascular disorders e.g. coronary arteriosclerosis and myocarditis; cancer e.g. melanoma and lymphoma. The proteins and polynucleotide sequences may also be used in wound healing and the treatment of infectious diseases. The human secreted protein gene #11 and protein sequences are represented in sequences AAA80616 and AAB25586. Sequences AAB25576-B25593. The human secreted proteins have various active dependent on the tissues in which they are expressed. Examples activities of the proteins include: immunosuppressant; The present invention relates to 12 secreted human proteins and the nucleotide sequences encoding them. The polynucleotide sequences given in AAA80606-A80623 encode the 12 secreted protein sequences given in sequences are represented AAA80677-A80682 represent genes related to the secreted protein activities gene#11

Sequence 229

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GCTATGGCCAGCAAGCCACCACGACTATCAGGTGGTTCAAAGGGAACACAGAGCTAAAA 447
                  Similarity:
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27-APR-2000;
20-JUN-2000;
19-JUL-2000;
01-SEP-2000;
15-SEP-2000;
20-OCT-2000;
                                                                                                                                                                                                                                                                                    Human; cytokine; cell proliferation; cell differentiation; gene therapy; vaccine; peptide therapy; stem cell growth factor; haematopolesis; tissue growth factor; immunomodulatory; cancer; leukaemia;
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2000US-0560875
2000US-0598075
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Examples; Fig
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This sequence is the human beta-secretase enzyme polynucleotide. Beta-secretase is capable of cleaving the beta-amyloid precursor prot (APP) (AN3742):wedish mutant APP). This enzyme is used in a method detecting human beta-secretase cleavage of polypeptides and for identifying beta-secretase inhibitors. The protein has a deduced molecular weight of 49 kD, although this is without glycosylation and the expected molecular weight is therefore higher. Inhibition of beta-secretase activity would be useful for chemical modelling of a critical event in the pathology of Alzheimer's disease. Inhibitors of beta-secretase would be useful for the prevention and treatment of Alzheimer's disease and Down's Syndrome. protein thod for of.

Sequence 444 AA

Alignment Scores:

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Percent Similarity:
Best Local Similarity:
Query Match:

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HisTyrThrProSerValLysIle-----IleProSerThrProPheProGlnGlu
                             CAGTATAAGCCTCAAGTGCACATTCAGATGACTTATCCTCTACAAGGCTTAACCCGGGAA
                                                        CAGGTGGAGCACCCTGCGGTCACTGGAAACCTGCAG---ACCCAGCGGTATCTAGAAGTA
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ArgValAspHisGluSerLeuAsnAlaThrProGlnValAlaMetGlnValLeuGluIle
                                                                                                                                         GTGACCAGTCAGCTGATGCTGAAGGTGCACAAGGAGGACGATGGGGTCCCAGTGATCTGC
                                                                                                                                                                                    {\tt LysGluIleLysAspValLysTyrLeuLysGluGluAspAlaAsnArgLysThrPheThr}
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Sequence 90, Appl Sequence 17, Appl Sequence 1133, Appl Sequence 80, Appl Sequence 89, Appl Sequence 67, Appl Sequence 71, Appl	23, 23, 24, 20, 20, 20, 20, 20, 20, 20, 20, 20, 20	10, 10, 10, 10, 10, 10, 10, 10, 10, 10,	20, Ap 22, Ap 22	sequence 6.

## ALIGNMENTS

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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P25488P1C1 CURRENT APPLICATION NUMBER: US/09/944,413 CURRENT FILING DATE: 2001-09-26 PRIOR APPLICATION NUMBER: 09/866,028 PRIOR FILING DATE: 2001-05-25
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                                                                                                                        Wood, William
                                                                                                                                        Roy, Margaret
Tumas, Daniel
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Napier, Mary
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Grimaldi, Christopher
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ICR APPLICATION NUMBER: 60/14, 296
ICR APPLICATION NUMBER: 60/146, 222
ICR FILING DATE: DECEmber 22, 1998
ICR APPLICATION NUMBER: 60/146, 222
ICR FILING DATE: September 16, 1998
ICR APPLICATION NUMBER: PCT/US98/19330
ICR FILING DATE: DECEmber 1, 1998
ICR APPLICATION NUMBER: PCT/US98/25108
ICR APPLICATION NUMBER: 09/216, 021
ICR FILING DATE: DECEmber 12, 1998
ICR APPLICATION NUMBER: 09/216, 021
ICR FILING DATE: DECEmber 22, 1998
ICR APPLICATION NUMBER: 09/218, 311
ICR FILING DATE: DECEmber 15, 1999
ICR APPLICATION NUMBER: PCT/US99/1252
ICR APPLICATION NUMBER: PCT/US99/21090
ICR FILING DATE: September 15, 1999
ICR APPLICATION NUMBER: PCT/US99/28301
ICR APPLICATION NUMBER: PCT/US99/28301
ICR FILING DATE: NO. US20020156004A1ember
ICR APPLICATION NUMBER: PCT/US99/28301
ICR APPLICATION NUMBER: PCT/US99/30095
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OR FILLING DATE: February 9, 1998
OR APPLICATION NUMBER: 60/075,945
OR FILLING DATE: February 25, 1998
OR APPLICATION NUMBER: 60/112,850
OR FILING DATE: December 16, 1998
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APPLICATION NUMBER: PCT/US00/32678 FILING DATE: December 1, 2000 APPLICATION NUMBER: PCT/US01/06520
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APPLICATION NUMBER: 60/069,873
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APPLICATION NUMBER: 60,709,197
APPLICATION NUMBER: 60,769,870
APPLICATION NUMBER: 60,769,870
December 17, 1997
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                                                             ATION NUMBER: PCT/US00/14042
DATE: May 22, 2000
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NUMBER OF SEQ ID NOS: 120
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CURRENT FILING DATE: 2001-09-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES TITLE OF INVENTION: ACIDS ENCODING THE SAME
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PRIOR APPLICATION NUMBER: 60/067,411
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APPLICATION NUMBER: 60/069,702
FILING DATE: December 16, 1997
APPLICATION NUMBER: 60/069,870
APPLICATION NUMBER: 17, 1997
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APPLICATION NUMBER: 60/069,278
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APPLICATION NUMBER: 60/074, 097
FILING DATE: February 9, 1998
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                                                                                                    FILING DATE: December 18, 199
APPLICATION NUMBER: 60/070, 44
FILING DATE: January 5, 1998
APPLICATION NUMBER: 60/074,08
FILING DATE: February 9, 1998
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APPLICATION NUMBER: 60/069,696
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APPLICATION NUMBER: 60/069,873
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OR APPLICATION NUMBER: 09/218 517
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OR APPLICATION NUMBER: 09/254,311
OR FILING DATE: March 3, 1999
OR APPLICATION NUMBER: PCT/US99/12252
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FILING DATE: March 2, 2000
APPLICATION NUMBER: PCT/US00/08439
FILING DATE: March 30, 2000
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APPLICATION NUMBER: PCT/US00/03565
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APPLICATION NUMBER: PCT/US00/04414
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                            Goddard, Audrey
Godowski, Paul
Grimaldi, Christopher
Gurney, Austin
Hillan, Kenneth
                                                                            Filvaroff, Ellen
Gerritsen, Mary
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                    Kljavin, Ivar
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           Napier,Mary
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PRIOR FILING DATE: MAY 22, 2000
PRIOR APPLICATION NUMBER: PCT/US00/20710
PRIOR FILING DATE: July 28, 2000
PRIOR APPLICATION NUMBER: PCT/US00/32678
PRIOR FILING DATE: December 1, 2000
PRIOR APPLICATION NUMBER: PCT/US01/06520
PRIOR APPLICATION NUMBER: PCT/US01/06520
PRIOR FILING DATE: February 28, 2001
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SEQ ID NO
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APPLICANT: Baker, Kevin
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                                                                                                                                                                                                                                                                                            APPLICANT: WOOD, WILLIAM
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P2548P1C1
CURRENT APPLICATION NUMBER: US/09/866,028
CURRENT APPLICATION NUMBER: US/09/866,028
CURRENT APPLICATION DATE: 2001-05-25
Prior application data removed - consult PALM o
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Hillan,Kenneth
Kljavin,Ivar
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Goddard, Audrey
Godowski, Paul
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Tumas, Daniel
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          GENERAL INFORMATION:

APPLICANT: Baker, Ke-
APPLICANT: Botstein

APPLICANT: Ferrara,

APPLICANT: Filvarof,

APPLICANT: Gerritse,

APPLICANT: Goddard,

APPLICANT: Goddwski
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US20020102647A1
Grimaldi, Christopher
             Goddard, Audrey
Godowski, Paul
                                                         Ferrara, Napoleone Filvaroff, Ellen
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OR APPLICATION NUMBER: 60/074 086
OR FILING DATE: February 9, 1998
OR APPLICATION NUMBER: 60/074 092
OR FILING DATE: February 9, 1998
OR APPLICATION NUMBER: 60/075, 945
OR FILING DATE: February 25, 1998
OR APPLICATION NUMBER: 60/112, 850
OR APPLICATION NUMBER: 60/113, 296
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 60/145, 296
OR FILING DATE: December 22, 1998
OR APPLICATION NUMBER: 60/146, 222
OR FILING DATE: July 28, 1999
OR APPLICATION NUMBER: FCT/US98/193:
OR APPLICATION NUMBER: FCT/US98/2510
OR APPLICATION NUMBER: PCT/US98/2510
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 09/216, 021
OR APPLICATION NUMBER: 09/218, 517
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OR FILING DATE: December 3, 199
OR APPLICATION NUMBER: 60/069,3
OR FILING DATE: December 11, 19
OR APPLICATION NUMBER: 60/069,3
OR FILING DATE: December 11, 19
OR APPLICATION NUMBER: 60/069,2
OR FILING DATE: December 12, 19
OR APPLICATION NUMBER: 60/069,6
OR FILING DATE: December 16, 19
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OR APPLICATION NUMBER: 60/069,7
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N NUMBER: 09/218,517

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                        NUMBER: PCT/US99/28301: December1, 1999
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PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: February 22, 2000
PRIOR PPLICATION NUMBER: PCT/US00/05841
PRIOR PPLICATION NUMBER: PCT/US00/05841
PRIOR FILING DATE: March 2, 2000
PRIOR PPLICATION NUMBER: PCT/US00/06439
PRIOR PRILING DATE: March 30, 2000
PRIOR APPLICATION NUMBER: PCT/US00/14042
PRIOR FILING DATE: May 22, 2000
PRIOR PILING DATE: July 28, 2000
PRIOR PILING DATE: July 28, 2000
PRIOR FILING DATE: December 1, 2000
PRIOR APPLICATION NUMBER: PCT/US00/32678
PRIOR APPLICATION NUMBER: PCT/US01/36520
PRIOR APPLICATION NUMBER: PCT/US01/36520
PRIOR PILING DATE: December 1, 2001
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PRIOR PILING DATE: February 28, 2001
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TITLE OF INVENTION: SECRETED AND TRANSMENTITLE OF INVENTION: ACIDS ENCODING THE STILL REFERENCE: P2548P1C1

CURRENT APPLICATION NUMBER: US/09/944,457

CURRENT APPLICATION NUMBER: 09/866,028

PRIOR APPLICATION NUMBER: 60/067,411

PRIOR APPLICATION NUMBER: 60/069,341

PRIOR APPLICATION NUMBER: 60/069,334

PRIOR FILING DATE: December 11, 1997

PRIOR APPLICATION NUMBER: 60/069,335

PRIOR FILING DATE: December 11, 1997

PRIOR APPLICATION NUMBER: 60/069,278

PRIOR FILING DATE: December 11, 1997

PRIOR APPLICATION NUMBER: 60/069,278

PRIOR FILING DATE: December 11, 1997

PRIOR APPLICATION NUMBER: 60/069,696

PRIOR FILING DATE: December 16, 1997

PRIOR APPLICATION NUMBER: 60/069,694

PRIOR APPLICATION NUMBER: 60/069,696

PRIOR APPLICATION NUMBER: 60/069,696

PRIOR APPLICATION NUMBER: 60/069,696

PRIOR FILING DATE: December 16, 1997

PRIOR APPLICATION NUMBER: 60/069,702

PRIOR FILING DATE: December 17, 1997

PRIOR APPLICATION NUMBER: 60/069,804
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Hillan, Kenneth
Kljavin, Ivar
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Godowski, Paul
Grimaldi, Christopher
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Tumas, Daniel
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OR APPLICATION NUMBER: 09/218,517
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 09/254,311
OR FILING DATE: March 3, 1999
OR APPLICATION NUMBER: 09/254,311
OR FILING DATE: March 3, 1999
OR APPLICATION NUMBER: PCT/US99/21252
OR FILING DATE: June 22, 1999
OR APPLICATION NUMBER: PCT/US99/21090
OR FILING DATE: September 15, 1999
OR APPLICATION NUMBER: PCT/US99/28409
OR FILING DATE: No. US20020110859Alember
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OR FILING DATE: December 17, 1997
OR APPLICATION NUMBER: 60/068,017
OR APPLICATION NUMBER: 60/070,440
OR FILING DATE: December 18, 1997
OR APPLICATION NUMBER: 60/070,440
OR FILING DATE: December 60/074,086
OR APPLICATION NUMBER: 60/074,086
OR APPLICATION NUMBER: 60/074,092
OR FILING DATE: February 9, 1998
OR APPLICATION NUMBER: 60/075,945
OR FILING DATE: February 25, 1998
OR APPLICATION NUMBER: 60/12,945
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 60/113,296
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APPLICATION NUMBER: PCT/US00/20710
FILING DATE: July 28, 2000
APPLICATION NUMBER: PCT/US00/32678
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NUMBER: PCT/US00/08439
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NUMBER: PCT/US00/14042
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: February 11, 2000
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PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
PRIOR APPLICATION NUMBER: 60/069,334
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES
TITLE OF INVENTION: ACIDS ENCODING THE SAME
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OR FILLING DATE: January 5, 1998
OR FILLING DATE: February 9, 1998
OR APPLICATION NUMBER: 60/074,092
OR FILLING DATE: February 9, 1998
OR APPLICATION NUMBER: 60/075,945
OR FILLING DATE: February 25, 1998
OR FILLING DATE: December 16, 1998
OR APPLICATION NUMBER: 60/112,850
OR FILLING DATE: December 16, 1998
OR APPLICATION NUMBER: 60/113,296
OR APPLICATION NUMBER: 60/113,296
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R FILING DATE: December 17, 1997
R APPLICATION NUMBER: 60/060,1997
R FILING DATE: December 18, 1997
PR APPLICATION NUMBER: 60/070,440
R FILING DATE: January 5, 1998
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ENT APPLICATION NUMBER: US/09/944,862
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FILING DATE: December 16, 1997
APPLICATION NUMBER: 60/069,694
FILING DATE: December 16, 1997
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APPLICATION NUMBER: 60/069,278
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Filvaroff, Ellen
Gerritsen, Mary
Goddard, Audrey
Goddwski, Paul
Grimaldi, Christopher
Gurney, Austin
Hillan, Kenneth
Kijavin, Ivar
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                                                                                                                                                                                                                                                            NUMBER: 60/146,222
: July 28, 1999
NUMBER: PCT/US98/19330
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NUMBER: 09/218,517
: December 22, 1998
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PRIOR FILING DATE: February 28, 2001
NUMBER OF SEQ ID NOS: 120
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APPLICATION NUMBER: PCT/US99/30095
FILING DATE: December 16, 1999
APPLICATION NUMBER: PCT/US00/03565
FILING DATE: February 11, 2000
APPLICATION NUMBER: PCT/US00/04414
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                                                                                                   ACAGTCCTGGTCCCACCACCACGTAATCTGATGATGATATCCAGAAAGACACTGCGGTGGAA
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                  TrpPheLysGlyAsnThrGluLeuLysGlyLysSerGluValGluGluTrpSerAspMet
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: NO. US20020115145A1ember 30, 1999
NUMBER: PCT/US99/28313
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                                                              CURRENT APPLICATION NUMBER: US/09/945, CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 09/866,028
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December 11, 1997
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APPLICATION NUMBER: 60/069335
FILING DATE: December 11, 1997
APPLICATION NUMBER: 60/069,278
FILING DATE: December 11, 1997
APPLICATION NUMBER: 60/069,425
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Tumas, Daniel
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OR APPLICATION NUMBER: 09/216,021
OR FILING DATE: December 16, 1998
IOR FILING DATE: December 22, 1998
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IOR APPLICATION NUMBER: 09/254,311
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OR FILING DATE: December: 60/11,396
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FILING DATE: MARCh 3, 1999
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FILING DATE: May 22, 2000
APPLICATION NUMBER: PCT/US00/20710
FILING DATE: July 28, 2000
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CURRENT APPLICATION NUMBER: US/09/945,015
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 09/866,028
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,328
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,425
PRIOR FILING DATE: December 16, 1997
PRIOR APPLICATION NUMBER: 60/069,628
PRIOR FILING DATE: December 16, 1997
PRIOR APPLICATION NUMBER: 60/069,696
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PRIOR APPLICATION NUMBER: 60/069,702
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PRIOR APPLICATION NUMBER: 60/069,873
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IOR APPLICATION NUMBER: 09/218,517
IOR FILING DATE: December 22, 1998
IOR APPLICATION NUMBER: 09/254,311
IOR FILING DATE: March 3, 1999
IOR APPLICATION NUMBER: PCT/US99/1252
IOR APPLICATION NUMBER: PCT/US99/1252
IOR APPLICATION NUMBER: PCT/US99/21090
IOR FILING DATE: Mo. US20020132768Alember 30
IOR APPLICATION NUMBER: PCT/US99/28301
IOR APPLICATION NUMBER: PCT/US99/28301
IOR FILING DATE: NO. US20020132768Alember 30
IOR APPLICATION NUMBER: PCT/US99/28301
IOR FILING DATE: December 1, 1999
IOR APPLICATION NUMBER: PCT/US99/30095
IOR FILING DATE: December 16, 1999
IOR APPLICATION NUMBER: PCT/US00/04414
IOR APPLICATION NUMBER: PCT/US00/04414
IOR FILING DATE: PEDRUARY 22, 2000
IOR APPLICATION NUMBER: PCT/US00/05841
IOR FILING DATE: PEDRUARY 2, 2000
IOR APPLICATION NUMBER: PCT/US00/05841
IOR FILING DATE: March 2, 2000
IOR APPLICATION NUMBER: PCT/US00/08439
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FILING DATE: February 28 200
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FILING DATE: May 22, 2000
APPLICATION NUMBER: PCT/US00/20710
FILING DATE: July 28, 2000
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                                                                              APPLICANT: Napier, Mary
APPLICANT: Roy, Margaret
APPLICANT: Tunas, Daniel
APPLICANT: Wood, William
TITLE OF INVENTION: SECRETED AN
TITLE OF INVENTION: ACIDS ENCE
FILE REFERENCE: P2548P1C1
                                                                                                                                                                                    APPLICANT Baker, A
APPLICANT BOTSTEI
APPLICANT FETTATE
APPLICANT FILVATOR
APPLICANT GODOWS
APPLICANT GOODWS
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                                     CURRENT FILING DATE: 2001-09-26
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APPLICATION NUMBER: 09/866,028 FILING DATE: 2001-05-25
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Godowski, Paul
Grimaldi, Christopher
Gurney, Austin
Hillan, Kenneth
Kljavin, Ivar
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RIOR FILING DATE: December 11, 1997

RIOR APPLICATION NUMBER: 60/069, 425

RIOR FILING DATE: December 12, 1997

RIOR APPLICATION NUMBER: 60/069, 425

RIOR FILING DATE: December 16, 1997

LIOR APPLICATION NUMBER: 60/069, 696

LIOR FILING DATE: December 16, 1997

LOR APPLICATION NUMBER: 60/069, 696

LOR FILING DATE: December 16, 1997

LOR APPLICATION NUMBER: 60/069, 702

LOR FILING DATE: December 17, 1997

LOR APPLICATION NUMBER: 60/069, 870

LOR FILING DATE: December 17, 1997

LOR APPLICATION NUMBER: 60/069, 870

LOR FILING DATE: December 17, 1997

LOR APPLICATION NUMBER: 60/069, 873

LOR APPLICATION NUMBER: 60/070, 440

LOR FILING DATE: December 17, 1998

LOR APPLICATION NUMBER: 60/074, 092

LOR APPLICATION NUMBER: 60/112, 296

LOR APPLICATION NUMBER: 60/113, 296

LOR APPLICATION
OR APPLICATION NUMBER: PCT/US99/12252

OR FILING DATE: June 22, 1999

OR APPLICATION NUMBER: PCT/US99/21090

OR FILING DATE: September 15, 1999

OR APPLICATION NUMBER: PCT/US99/28090

OR APPLICATION NUMBER: PCT/US99/2814

OR APPLICATION NUMBER: PCT/US99/28313

OR FILING DATE: No. US20020132981A1ember 30

OR APPLICATION NUMBER: PCT/US99/28301

OR APPLICATION NUMBER: PCT/US99/30095

OR APPLICATION NUMBER: PCT/US99/30095

OR APPLICATION NUMBER: PCT/US99/30565

OR FILING DATE: December 16, 1999

OR APPLICATION NUMBER: PCT/US00/03565

OR APPLICATION NUMBER: PCT/US00/05841

OR APPLICATION NUMBER: PCT/US00/05841

OR APPLICATION NUMBER: PCT/US00/06439

OR APPLICATION NUMBER: PCT/US00/04439

OR APPLICATION NUMBER: PCT/US00/14042

OR APPLICATION NUMBER: PCT/US00/20710

OR APPLICATION NUMBER: PCT/US00/20710

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R APPLICATION NUMBER: 09/216,021
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NUMBER OF SEQ ID NOS:
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES.
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P2548P1C1
CURRENT APPLICATION NUMBER: U5/09/944,097
CURRENT FILING DATE: 2001-08-31
PRIOR APPLICATION NUMBER: 09/866,028
PRIOR FILING DATE: 2001-05-25
PRIOR FILING DATE: 2001-05-25
PRIOR FILING DATE: December 11, 1997
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OR FILING DATE: December 11, 1997
OR APPLICATION NUMBER: 60/069, 425
OR FILING DATE: December 12, 1997
OR APPLICATION NUMBER: 60/069, 696
OR FILING DATE: December 16, 1997
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                                                                  APPLICATION NUMBER: 60/074,092
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                                  FILING DATE: February 25, 1 APPLICATION NUMBER: 60/112,
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APPLICATION NUMBER: PCT/US00/20710
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FILING DATE: December 16, 1999
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APPLICATION NUMBER: 09/254,311
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APPLICANT:
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Goddard, Audrey
Godowski, Paul
Grimaldi, Christopher
Gurney, Austin
Hillan, Kenneth
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Napler, Mary
Roy, Margaret
Tumas, Daniel
Wood, William
                                                                                 Eaton, Dan
Ferrara, Napoleone
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                PRIOR
      IOR APPLICATION NUMBER: 60/067,411
IOR FILING DATE: December 3, 1997
IOR APPLICATION NUMBER: 60/069,334
IOR FILING DATE: December 11, 1997
IOR APPLICATION NUMBER: 60/069,278
IOR FILING DATE: December 11, 1997
IOR APPLICATION NUMBER: 60/069,278
IOR FILING DATE: December 11, 1997
IOR APPLICATION NUMBER: 60/069,694
IOR FILING DATE: December 16, 1997
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IOR APPLICATION NUMBER: 60/069,870
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IOR APPLICATION NUMBER: 60/113,896
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IOR APPLICATION NUMBER: PCT/US98/25138
IOR APPLICATION NUMBER: PCT/US99/281313
IOR FILING DATE: December 16, 1998
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OR FILING DATE: May 22, 2000
OR APPLICATION NUMBER: PCT/US00/20710
OR APPLICATION NUMBER: PCT/US00/32678
OR APPLICATION NUMBER: PCT/US00/32678
OR APPLICATION NUMBER: PCT/US01/06520
OR APPLICATION NUMBER: PCT/US01/06520
OR FILING DATE: February 28, 2001
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APPLICANT: Bater, Kevin
APPLICANT: Bater, Kevin
APPLICANT: Botstein, David
APPLICANT: Botstein, David
APPLICANT: Firrara, Napoleone
APPLICANT: Firraraff, Ellen
APPLICANT: Firraraff, Ellen
APPLICANT: Gerittsen, Mary
APPLICANT: Gerittsen, Mary
APPLICANT: Gerittsen, Mary
APPLICANT: Gerittsen, Mary
APPLICANT: Grimaldi, Christopher
APPLICANT: Wood, William
APPLICANTON NUMBER: US/09/943,762
CURRENT APPLICATION NUMBER: US/09/943,762
CURRENT APPLICATION NUMBER: 00/067,411
PRIOR FILING DATE: December 13, 1997
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,335
PRIOR APPLICATION NUMBER: 60/069,788
PRIOR APPLICATION NUMBER: 60/069,425
PRIOR APPLICATION NUMBER: 60/069,694
PRIOR FILING DATE: December 12, 1997
PRIOR APPLICATION NUMBER: 60/069,894
PRIOR APPLICATION NUMBER: 60/069,894
PRIOR FILING DATE: December 15, 1997
PRIOR APPLICATION NUMBER: 60/069,894
PRIOR FILING DATE: December 16, 1997
PRIOR APPLICATION NUMBER: 60/069,894
PRIOR FILING DATE: December 16, 1997
PRIOR APPLICATION NUMBER: 60/069,894
PRIOR FILING DATE: December 16, 1997
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PRIOR FILING DATE: December 16, 1997
PRIOR APPLICATION NUMBER: 60/069,894
PRIOR APPLICATION NUMBER: 60/069,89
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IOR APPLICATION NUMBER: 60/113,296
IOR APPLICATION NUMBER: 60/146,222
IOR FILING DATE: December 22, 1998
IOR APPLICATION NUMBER: 60/146,222
IOR FILING DATE: September 16, 1998
IOR APPLICATION NUMBER: PCT/US98/1930
IOR APPLICATION NUMBER: PCT/US98/25108
IOR APPLICATION NUMBER: PCT/US98/25108
IOR APPLICATION NUMBER: PCT/US98/25108
IOR APPLICATION NUMBER: 09/216,021
IOR FILING DATE: December 16, 1998
IOR APPLICATION NUMBER: 09/254,311
IOR FILING DATE: December 22, 1998
IOR APPLICATION NUMBER: PCT/US99/1252
IOR FILING DATE: March 3, 1999
IOR APPLICATION NUMBER: PCT/US99/21090
IOR APPLICATION NUMBER: PCT/US99/21090
IOR APPLICATION NUMBER: PCT/US99/28409
IOR APPLICATION NUMBER: PCT/US99/28409
IOR APPLICATION NUMBER: PCT/US99/28301
IOR APPLICATION NUMBER: PCT/US99/28301
IOR APPLICATION NUMBER: PCT/US99/28301
IOR APPLICATION NUMBER: PCT/US99/28301
IOR APPLICATION NUMBER: PCT/US99/30095
IOR APPLICATION NUMBER: PCT/US99/30095
IOR APPLICATION NUMBER: PCT/US99/30095
IOR APPLICATION NUMBER: PCT/US90/05641
IOR APPLICATION NUMBER: PCT/US00/05641
IOR APPLICATION NUMBER: PCT/US00/05641
IOR APPLICATION NUMBER: PCT/US00/05641
IOR APPLICATION NUMBER: PCT/US00/06439
IOR APPLICATION NUMBER: PCT/US00/14042
IOR APPLICATION NUMBER: PCT/US00/14042
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R APPLICATION NUMBER: PCT/US01/06520
R FILING DATE: February 28, 2001
R OF SEQ ID NOS: 120
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                                                                                                  N NUMBER: 60/075 945

FE: February 25, 1998

N NUMBER: 60/113, 850

FE: December 16, 1998

N NUMBER: 60/113, 296

NE: December 22, 1998

NN NUMBER: 60/146, 222

FE: July 28, 1999
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Sequence 61, Application US/
Patent No. US20020142959A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin
APPLICANT: Botstein, David
APPLICANT: Eaton, Dan
APPLICANT: Ferrara, Napolec
APPLICANT: Filvaroff, Eller
APPLICANT: Gerritsen, Mary
APPLICANT: Goddard, Audrey
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                                                 TTCATCAATAACCTAAACAAAACAGATAATGGTACATACCGCTGTGAAGCTTCAAACATA
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Eaton, Dan Ferrara, Napoleone Filvaroff, Ellen Gerritsen, Mary Botstein, David

US/09944654

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CURRENT APPLICATION NUMBER: US/09/944,654
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 09/866,028
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
PRIOR FILING DATE: December 11, 1997
PRIOR FILING DATE: December 11, 1997
PRIOR FILING DATE: December 11, 1997
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APPLICANT:
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IOR FILING DATE: February 9, 1998
IOR APPLICATION NUMBER: 60/075,945
IOR FILING DATE: February 25, 1998
IOR APPLICATION NUMBER: 60/112,850
IOR FILING DATE: December 16, 1998
IOR APPLICATION NUMBER: 60/113,296
IOR FILING DATE: December 22, 1998
IOR APPLICATION NUMBER: 60/146,222
IOR FILING DATE: December 22, 1998
IOR APPLICATION NUMBER: 60/146,222
IOR FILING DATE: September 16, 1998
IOR APPLICATION NUMBER: PCT/US98/19330
IOR APPLICATION NUMBER: 09/216,021
IOR FILING DATE: December 16, 1998
IOR APPLICATION NUMBER: 09/216,021
IOR APPLICATION NUMBER: 09/218,517
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OR APPLICATION NUMBER: 60/669,870
OR FILING DATE: December 17, 1997
OR APPLICATION NUMBER: 60/669,873
OR FILING DATE: December 17, 1997
OR APPLICATION NUMBER: 60/068,017
OR FILING DATE: December 18, 1997
OR APPLICATION NUMBER: 60/070,440
OR FILING DATE: January 5, 1988
OR APPLICATION NUMBER: 60/074,086
OR FILING DATE: February 9, 1998
OR FILING DATE: February 9, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OR APPLICATION NUMBER: 60/069,696
OR FILLING DATE: December 16, 1997
OR APPLICATION NUMBER: 60/069,694
OR FILLING DATE: December 16, 1997
OR APPLICATION NUMBER: 60/069,702
OR FILLING DATE: December 16, 1997
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APPLICATION NUMBER: PCT/US99/21090
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APPLICATION NUMBER: 60/069,425
FILING DATE: December 12, 1997
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FILING DATE: December 11, 199
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Grimaldi, Christ
Gurney, Austin
Hillan, Kenneth
Kljavin, Ivar
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Roy, Margaret
Tumas, Daniel
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December 22, 1998
N UMBER: 09/254,311
E: MAICH 3, 1999
N UMBER: 09/254,311
                                                NUMBER: PCT/US99/28409

NO. US20020142959Alember

NUMBER: PCT/US99/28313

NO. US20020142959Alember
                                                                                                                                                           NUMBER: PCT/US99/12252
                                     NUMBER: PCT/US99/28301
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Best Local Similarity:
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PRIOR FILING DATE: March 2, 2000
PRIOR APPLICATION NUMBER: PCT/US00/08439
PRIOR FILING DATE: March 30, 2000
PRIOR APPLICATION NUMBER: PCT/US00/14042
PRIOR APPLICATION NUMBER: PCT/US00/20710
PRIOR APPLICATION NUMBER: PCT/US00/20710
PRIOR APPLICATION NUMBER: PCT/US00/32678
PRIOR APPLICATION NUMBER: PCT/US00/32678
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FILING DATE: February 11, 2000
APPLICATION NUMBER: PCT/US00/04414
FILING DATE: February 22, 2000
FILING DATE:
                                                                                                 TACACTGTGACCAGTCAGCTGATGCTGAAGGTGCACAAGGAGGACGATGGGGTCCCAGTG
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P2548P1C1
CURRENT APPLICATION NUMBER: US/09/943,851A
CURRENT APPLICATION NUMBER: US/09/966,028
PRIOR APPLICATION NUMBER: US/09/866,028
PRIOR FILING DATE: 2001-08-30
PRIOR FILING DATE: DOcember 3, 1997
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 1, 1997
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,335
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,278
PRIOR APPLICATION NUMBER: 60/069,425
PRIOR APPLICATION NUMBER: 60/069,425
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,696
PRIOR FILING DATE: December 16, 1997
PRIOR APPLICATION NUMBER: 60/069,696
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PRIOR FILING DATE: December 16, 1997
PRIOR APPLICATION NUMBER: 60/069,709
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US-09-943-851A-61
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Goddard, Audrey
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PRIOR APPLICATION NUMBER: 60/112,850
PRIOR APPLICATION NUMBER: 60/113,296
PRIOR FILING DATE: December 16, 1998
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RIOR APPLICATION NUMBER: 60/146,222
21OR FILING DATE: July 28, 1999
TOR APPLICATION NUMBER: PCT/US98/19330
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IOR FILING DATE: December 16, 1998
IOR APPLICATION NUMBER: 09/218,517
IOR FILING DATE: December 22, 1998
IOR APPLICATION NUMBER: 09/254,311
IOR FILING DATE: March 3, 1999
IOR APPLICATION NUMBER: PCT/US99/1225
IOR FILING DATE: June 22, 1999
IOR APPLICATION NUMBER: PCT/US99/281090
IOR APPLICATION NUMBER: PCT/US99/28409
IOR APPLICATION NUMBER: PCT/US99/28301
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APPLICATION NUMBER: PCT/US00/00414
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APPLICATION NUMBER: PCT/US00/05841
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FILING DATE: February 9, 1998
APPLICATION NUMBER: 60/074,092
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Sequence 12, Appl	Sequence 12, Appl	Sequence 12, Appl	Sequence 62, Appl	Sequence 21, Appl	Sequence 19, Appl	15	8	Sequence 60, Appl	Sequence 33, Appl	Sequence 29, Appl	Sequence 16, Appl	Sequence 2, Appli	Sequence 25, Appl	Sequence 17, Appl	Sequence 3, Appli		Sequence 45, Appl

ALIGNMENTS

#### RESULT 1 US-08-659-984A-1 Sequence 1, Application US/08659984A Patent No. 5942400 GENERAL INFORMATION: PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/485,152 FILING DATE: 07-JUN-1995 ATTORNEY/AGENT INFORMATION: NAME: Heslin, James M. REGISTRATION NUMBER: 29,541 REFERENCE/DOCKET NUMBER: 15270-0028: TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: TELECOMMUNICATION 115-326-2400 TELEFAX: 415-326-2422 INFORMATION FOR SEQ ID NO: 1: ZIP: 94111-3834 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICANT: Anderson, John P. APPLICANT: Sinha, Sukanto APPLICANT: Jacobson-Croak, Kirsten L. TITLE OF INVENTION: Assays for Detecting Beta-Secretase TITLE OF INVENTION: Inhibition TOPOLOGY: 11 MOLECULE TYPE: NUMBER OF SEQUENCES: 2 CORRESPONDENCE ADDRESS: SEQUENCE CHARACTERISTICS: APPLICATION NUMBER: US/08/659,984A FILING DATE: 07-JUN-1996 CLASSIFICATION: 436 ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Ctr., 8th Floor CITY: San Francisco STATE: California COUNTRY: USA TYPE: amino acid STRANDEDNESS: LENGTH: 421 amino acids linear protein single 15270-002810US

Query Match Best Local Similarity

35.4%; 66;

Score 623.5; DB 2 Pred. No. 1.1e-46;

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Mismatches

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                                                                 MOLECULE TYPE:
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PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/480,498
FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
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                                                                                                STRANDEDNESS:
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                                                                                                                                                                                                                                                   NAME: Heslin, James M. REGISTRATION NUMBER: 2
                                                                                   TOPOLOGY:
                                                                                                                     TYPE:
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INVENTION: Beta-Secretase
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                                                                                                                                                                                                                           CLASSIFICATION: 436
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08,
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Heslin, James M.
                                                                                                  TELEFAX: 415-326-2422
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: 07-JUN-199
                  MOLECULE TYPE:
                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: 415-326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Assays for Detecting Beta-Secretase TITLE OF INVENTION: Inhibition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          307 LPTTIIPSLTTATVTTTVAITTSPTTSATTSSIRDPNALAGQNGP----DH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 242 WVRVDDEM--PQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTI 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    190
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: TWO ______CITY: San Francisco
                              STRANDEDNESS:
TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   183 QVEHPAVTGNLQ-TQRYLEVQYKPQVHIQMTYPLQGLTREGDALELTCEAIGKPQPVMVT 241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      126 VNCTAMASKPATTIRWFKGNTELKGKSEVEEWS---DMYTVTSQLMLKVHKEDDGVPVIC
                                                                     TYPE:
                                                                                                                                                                                                           NAME: Heslin, James M. REGISTRATION NUMBER: :
                                                                                   LENGTH:
                                                                                                                                                                                         REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          70
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RVDHESLNATPQVAMQVLEIHYTPSVKI~--IPSTPFPQEGQPLILTCESKGKPLPEPVL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LTCKTSGSKPAADIRWFKNDKEIKDVKYLKEEDANRKTFTVSSTLDFRVDRSDDGVAVIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WHELSISVSDVSLSDEGQYTCSLFTMPVKTSKAYLTVLGVPEKPQISGFSSPVMEGDLMQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SSELKVSLINVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVEGEEIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GQFPLTQNVTVVEGGTAILTCRVDQNDNTSLQWSNPAQQTLYFDDKKALRDNRIELVRAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GQNLFTKDVTVIEGEVATISCQVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQLLNFS
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                                                                   amino acid
                                                                                 444 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Two Embarcadero Ctr., 8th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sinha, Sukanto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anderson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                    IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                protein
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                                             single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Crew LLF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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Wed Nov 20 08:04:02 2002

4

GenCore version 5.1.3 Copyright (c) 1993 - 2002 Compugen Ltd.

OM protein - protein search, using sw model

Run on: November 20, 2002, 07:34:43; Search time 15 Seconds (without alignments) 2153.411 Million cell updates/sec

Title: Perfect score: Sequence:

US-09-778-187B-2\_COPY\_39\_374
1761
1 IPTGDGQNLFTKDVTVIEGE.....LTIITDSRAGEEGSTRAVDH 336

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched: 283224 segs, 96134422 residues

Total number of hits satisfying chosen parameters: 283224

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database : PIR\_73:\*
1: pir1:\*
2: pir2:\*
3: pir3:\*
4: pir4:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	8	7	6	ر. د	4	ω	N	1	NO.	Result
202.5	203.5	206	208.5	209.5	210	210.5	213.5	213.5	214	216	218	218		221	221	221	222	222	223	223	223.5	224	227	239	243	248	263.5	263.5	Score	1
11.5	11.6	11.7	11.8	11.9	11.9	12.0	12.1	12.1	12.2	12.3	12.4	12.4	12.4	12.5	12.5	12.5	12.6	12.6	12.7	12.7	12.7	12.7	12.9	13.6	13.8	14.1	15.0	15.0	Match	
345	765	702	345	584	1011	345	932	812	4391	478	417	392	518	538	417	392	467	344	1092	725	7962	530	4162	1088	725	407	5198	5175	Length	
2	N	N	N	N	N	N	N	2	N	N	ب	Н	N	N	N	N	ب	N	<u></u> -	N	N	N	N	ب	N	N	N	N	BG	
JC1239	C42632	A36319	JC4025	I50419	T13669	S03199	A42632	B42632	A38096	I53960	RWHUPA	RWHUPD	JC4024	168093	A44194	B44194	HLMSP3	I56551	JN0635	JE0100	I38346	A53437	T42633	IJXLNL	JE0099	T08732	T43290	T20992	ID	
	cell adhesion mole	carcinoembryonic a	opioid-binding cel	s-gicerin precurso		opioid-binding pro	cell adhesion mole	cell adhesion mole	rec			poliovirus recepto	poliovirus recepto	PRR2 delta - human		poliovirus recepto	poliovirus recepto	_		neural cell adhesi	ct.	poliovirus recepto	connectin/titin -	cell	D)	hypothetical prote	Ř.	hypothetical prote	Description	

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231 AIGKPQPVMVTWVRVDDEMPQHAVL----SGPNLFINNLNKTDNGTYRCEASNIVGKAHS 286

44 45	43	4 1	40	39	38	37	36	35	34	S S	32	31	30
189.5 189	191.5	193	193	194	195	195.5	197.5	200	201.5	202	202	202.5	202.5
10.8	10.9	11.0	11.0	11.0	11.1	11.1	11.2	11.4	11.4	11.5	11.5	11.5	11.5
646 1036	976	1344	761	338	1612	1091	3707	1323	338	1443	847	868	862
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138049 S22383	T29583	T14316	IJHUNG	JC5519	T30805	IJCHNL	S18252	PN0568	JC1238	I50600	JH0371	A46512	I49583
cell surface glyco axonin 1 precursor	hypothetical prote	rig-1 protein - mo	neural cell adhesi	50K glycoprotein p	duttl protein - mo	neural cell adhesi	heparan sulfate pr	connectin 3B - chi	opioid-binding pro	neogenin - chicken	B-cell adhesion pr	CD22 homolog/B lym	differentiation an

## ALIGNMENTS

Qy Db	Qy Db	Qy Db	Qy Db		1 · A A A	2222	AAAAS	<b>8 9 9 9</b>	2 2 2 2 S	80000B	R
Y 174 EDDGVPVICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDALELTCE 230	Y 128 CTAMASKPATTIRWFKGNTELKGKSEVEEWSDMYTVTSQLMLKVHK 173 	Y 70 KVSLTNVSISDEGRYFCQLYTDPPQESYT-TITVLVPPRNLMIDIQKD-TAVEGEEIEVN 127 ::::  ::  :  :  :  :  :  :  :  :  :  :	Y 14 VTVIBGEVATISCQVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQLLNFSSSEL 69 	Query Match 15.0%; Score 263.5; DB 2; Length 5175; Best Local Similarity 24.6%; Pred. No. 6.2e-10; Matches 87; Conservative 66; Mismatches 129; Indels 71; Gaps 15;	A;Gene: CESP:F15G9.4a A;Gene: CESP:F15G9.4a A;Map position: X A;Introns: 85/1; 120/1; 334/3; 370/1; 477/2; 606/3; 664/1; 935/3; 977/1; 1051/3; 1184 A;Introns: 85/1; 120/1; 334/3; 370/1; 477/2; 606/3; 664/1; 935/3; 977/1; 1051/3; 1184 ; 2512/2; 2593/3; 2699/3; 2759/1; 2852/1; 2889/3; 2913/3; 2941/1; 2967/3; 2991/3; 303 1; 4225/1; 4361/1; 4408/1; 4456/1; 4498/1; 4647/3; 4838/1; 4879/1; 4941/1; 5011/1; 50	A;Residues: 1-5175 <wi2> A;Residues: 1-5175 <wi2> A;Cross-references: EMBL:Z47070; PIDN:CAA87.344.1; GSPDB:GN00028; CESP:F15G9.4a A;Experimental source: clone T09B9</wi2></wi2>	submitted to the EMBL Data Library, December 1994 A;Reference number: Z19929 A;Accession: T24733 A;Status: preliminary; translated from GB/EMBL/DDBJ A;Molecule type: DNA	A;Residues: 1-5175 <wil> A;Residues: 1-5175 <wil> A;Residues: 1-5175 <wil> A;Cross-references: EMBL:Z47068; PIDN:CAA87335.1; GSPDB:GN00028; CESP:F15G9.4a A;Experimental source: clone F15G9 R;Korshaw, J.</wil></wil></wil>	submitted to the EMBL Data Library, December 1994 A;Reference number: Z19355 A;Accession: T20992 A;Status: preliminary; translated from GB/EMBL/DDBJ A;Wolecule type: DNA	cal protein F15G9.4a - Cae: Caenorhabditis elegans 5-Oct-1999 #sequence_revis on: T20992; T24733	RESULT 1

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2468

DEKVSMLVAPSFDEPNIVRRITVNSGNPSTLHCPAKGSPSPTITWLKDGNAIE

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C; Species: Ca
C; Date: 11-Ja
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  Qγ
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C;Genetics:
A;Gene: him-4; F15G9.4b
A;Gene: him-4; F15G9.4b
A;Introns: 85/1; 120/1; 334/3; 370/1; 477/2; 606/3; 664/1; 935/3; 977/1; 1051/3; 1184/3;
A;Introns: 85/1; 120/1; 334/3; 370/1; 477/2; 606/3; 2913/3; 2941/1; 2967/3; 2991/3; 3033/2; 2512/2; 2593/3; 2699/3; 2759/1; 2852/1; 2889/3; 2913/3; 2941/1; 2967/3; 2991/3; 3033/2; 4225/1; 4361/1; 4408/1; 4456/1; 4498/1; 4647/3; 4838/1; 4902/1; 4964/1; 5034/1; 5100/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Residues: 1-5198 <WIL>
A; Cross references: EMBL.247068; PIDN:CAA87336.1;
A; Experimental source: clone F15G9
R; Kershaw, J.
submitted to the EMBL Data Library, December 1994
A; Reference number: Z19929
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Cross-references: EMBL:AF074901; R;Sulston, J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R; Vogel, B.E.; Hedgecock, E.M. submitted to the EMBL Data Library, J. A; Description: Hemicentin is required
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            δÃ
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A; Residues: 1-5198 <WI2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Reference number: A; Accession: T24734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: mRNA
A; Residues: 1-5198 < VOG>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Status: preliminary; translated from GB/EMBL/DDBJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Status: preliminary; translated from GB/EMBL/DDBJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Reference number: Z22396
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                                             2409
                                                                                                                                                                                                                                                                                                                                                                             2200 VTAIKGGALPFKCPID--DDK----NFKGQIIWLRNYQPIDLEAEDARITRL---SNDR
    287
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                                           VYARP-PAKITWLKAGKPLQSDKFVKTSANGQKLYLFKLRETDSSKYTCIATNEAGTDKR
                                                                                                                                                                                                                                                                                                                      KVSLTNVSISDEGRYFCQLYTDPPQESYT-TITVLVPPRNLMIDIQKD-TAVEGEEIEVN 127
                                                                                                                                                                                                                                                                                                                                                                                                                      VTVIEGEVATISCQVNKSDDSVIQLLNPNRQTIYFRDFRPL----KDSRFQLLNFSSSEL
DYMLYVYDPPTTIPP
                                                                                                                                                                                                             CPA-TGKPEPDITWFKDGEAIHIENIADIIPNGELNG---
                                                                                                                                                                                                                                                     CTAMASKPATTIRWFKG---
                                                                                                                                                                                                                                                                                              RLTILNVTENDEGQYSCRVKNDAGENSFDFKATVLVPPTIIMLDKDKNKTAVEHSTVTLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        VYARP-PAKITWLKAGKPLQSDKFVKTSANGQKLYLFKLRETDSSKYTCIATNEAGTDKR
                                                                              AIGKPOPVMVTWVRVDDEMPQHAVL----SGPNLFINNLNKTDNGTYRCEASNIVGKAHS
                                                                                                                            EGDAGKYTCEADNSA - - GSVEQDVNVNVITIPKIEKDGIPSDYESQ - -
                                                                                                                                                                    EDDGVPVICQVEHPAVTGNLQTQRYLEVQYKPQVH----IQMTYPLQGLTREGDALELTCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DFKVSMLVAPSFDEPNIVRRITVNSGNPSTLHCPAKGSPSPTITWLKDGNAIE 2520
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DYMLYVYDPPTTIPP
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Caenorhabditis elegans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  87;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15.0%;
24.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              66;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PIDN:CAA87345.1; GSPDB:GN00028;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 263.5; DB 2
Pred. No. 6.2e-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. 6.26
5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PIDN: AAC26792.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    December
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         June
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                129;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GSPDB:GN00028;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       #text_change 18-Feb-2000
                                                                                                                                                                                                           -----NQLKITRIK 2354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 5198;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mediated
                                                                                                                            --QNERVVISCP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                71;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CESP:F15G9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CESP: F15G9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cell
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    germ
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JE0099
JE0099
neural cell adhesion molecule 1 - African clawed trog
neural cell adhesion molecule 1 - African clawed trog
N;Alternate names: N-CAM 1
C;Species: Xenopus laevis (African clawed frog)
C;Species: Xenopus laevis (African clawed frog)
The results of t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Kudo, M.; Takayama, E.; Tadakuma, T.; Shlokawa, K. Biochem. Biophys. Res. Commun. 245, T27-132, 1998 A;Title: Molecular cloning of s245, T27-132, 1998 A;Reference number: JE0099; MUID:98204770; PMID:9535795
                                 δõ
                                                                                                                                                                                                                                                                                                                                                                                                                       A; Molecule type: mRNA
A; Residues: 1-725 < KUD>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Reference number: A; Accession: JE0099
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Date: 19-May-1998 #sequence_revision
C;Accession: JE0099
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Вþ
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A;Experimental source: fetal kidney; clone DKF2p566B0846
C;Genetics:
A;Note: DKF2p566B0846.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Reference number: Z16474
A; Accession: T08732
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hypothetical protein DKFZp566B0846.1 - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change
                                                                                                                                                                                                                                                                                                                                                                               A;Cross-references: DDBJ:AB008162; NID:g3116226; PIDN:BAA25931.1; PID:g3116227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Residues: 1-407 <OTT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              submitted to the Protein Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T08732
                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       233 ADIEDLATEPKKLPFPLSTLATIKDDTIATIIASVVGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               138 TIRWFKGNTELKGKSEVEEWSDMY-----TVTSQLMLKVHKEDDGVPVICQVEHPAVTGN 192
                                                                                                                             Local
                          ,12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61 HIDW-EGDL-----GEMESTTTSFPNETATIISQYKLFPTRFARGRRITCVVKHPALEKD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  82 GRYFCQLYTDP--PQESYTTITVLVPPRNLMIDIQKDTAVEG--EEIEVNCTAMASKPAT 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 GKYICKAVTFPLGNAQSSTTVTVLVEPTVSLIK-GPDSLIDGGNETVAAICIAATGKPVA 60
LLASDNTLHFVHPLTFNYSGVYICKVTNSLGQRSDQKVIYISDPPTTTTLQPTIQWHPST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AVLSGPNL-FINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPP--TTIPP------
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LQTQRYLEVQYKPQVHIQMTYPLQGLTREGDALELTCEAIGKPQPVMVTWVRVDDEMPQH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IRYSFILDIQYAPEVSVTGYDGNWFVGRKG--VNLKCNADANPPPFKSVWSRLDGQWPDG
                                                                                                   90; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           77; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            B.; Obermaier,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mRNA
                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -PTTTTTTTTTTTLTILTITDSRAG
                                                                                                                                 13.8%;
26.5%;
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27.7%;
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                                                                                                   61;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        44;
                                                                                               Score 243; DB Pred. No. 1.5e 51; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 248; DB 2;
Pred. No. 3.3e-10,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Database,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mewes, H.W.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .W.; Gassenhuber, J.;
May 1999
                                                                                                   .5e-09;
les 147;
                                                                                                                                                                  DB
                                                                                                                                                                  2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           115;
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                                                                                                                                                              Length
                                                                                                      Indels
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                                                                                                      42;
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                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  232
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    as
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GenCore version 5.1.3 Copyright (c) 1993 - 2002 Compugen Ltd.

OM protein - protein search, using sw model

Run on: November 20, 2002, 07:34:42; Search time 9 Seconds (without alignments) 1548.450 Million cell updates/sec

Title: Perfect score: Sequence:

US-09-778-187B-2\_COPY\_39\_374
1761
1 IPTGDGQNLFTKDVTVIEGE.....LTIITDSRAGEEGSIRAVDH 336

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched: 112892 seqs, 41476328 residues

Total number of hits satisfying chosen parameters: 112892

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

SwissProt\_40:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

33	ب د د	ب د د	2 6	200	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	60	7	6	5	4	ω	2	1	:::::	N O	Result
193	101	101	195.5	195.5	197.5	202	202	202.5	202.5	202.5	203	206	207	208	208	208.5	210	210.5	213	214.5	217	219	221	221	222	223	223.5	224	227.5	¥	239		Score	
			1 -										٠	•	•	•	•	•	12.1	•	•	•	12.5	•	•	•	•	•	12.9	•	13.6	٠	_	Query
848	7 6	ى د د د د د	1234	1091	3707	1443	1377	862	353	345	1461	702	847	1493	515	345	1242	345	4393	837	417	337	538	417	344	1092	837	530	517	515	1088		Length	
<b>-</b>	_ <sub>_</sub>	۰,	ــر د		_	1	<u>, , , , , , , , , , , , , , , , , , , </u>	_	_	<u> </u>	μ.	μ	$\vdash$	μ	H	-	_	<u>,                                    </u>	_	Н	_		ب	1	<del>ب</del>	<u>_</u>	<b>,</b>	_	۳	_	<u>, , , , , , , , , , , , , , , , , , , </u>	•	BB	
NCA1_HUMAN	NOAS BUMAN	CTOOL MOORE	RAT	NCA1_CHICK	PGBM_MOUSE	NEO1_CHICK	NEO1_RAT	CD22_MOUSE	CEPU_CHICK	OPCM_RAT	NEO1_HUMAN		CD22_HUMAN	NEO1_MOUSE	PVR1_MOUSE	OPCM_HUMAN	NPHN_MOUSE	OPCM_BOVIN			PVR_HUMAN	G55A_CHICK	PVR2_HUMAN	PVR_CERAE	NTRI_RAT	NCA2_XENLA	NCM2_MOUSE	PVR2_MOUSE	PVR1_HUMAN	PVR1_PIG	NCA1_XENLA		ID	
	9011	משון שוווע	ratt	gal	mus mus	Q90610 gallus gall	P97603 rattus norv	329 mus mu:	3 gallus	736 rattus	59 homo	6731 homo	3 homo	98 mus	5 mus π	2 homo	7 mus	t sod t		4 homo	l homo s	2 gallu	N	5 cercopi	3 rattus n	5 xeno	sum s	7 mus r	homo sap	5 sus scro	P16170 xenopus lae		Description	

ပ်	44	43	42	41	40	9	38	37	3 6	35	34
182.5	183.5	185.5	187	187.5	188	188	188.5	189	189.5	190	192.5
10.4	10.4	10.5	10.6	10.6	10.7	10.7	10.7	10.7	10.8	10.8	10.9
873	338	338	588	583	858	853	764	1036	646	1447	1241
L		_		L	۳	_	-	H	μ	μ	Ľ
FAS2_DROME	LAMP_RAT	LAMP_HUMAN	C166_CHICK	C166_HUMAN	NCA1_RAT	NCA1_BOVIN	ICCR_DROME	AXO1_CHICK	MU18_HUMAN	DCC_MOUSE	NPHN_HUMAN
P34082	Q62813	Q13449	P42292	Q13740	P13596	P31836	008180	P28685	P43121	P70211	060500

# ALIGNMENTS

DR D	388888888	3888888	36666666	RRRRRROO	OC OC GE	AC DT DT DT	RESULT NCA1_X ID N
•	his SWISS-PROT entry is copyright. It is produced through a collaboratice tween the Swiss Institute of Bioinformatics and the EMBL outstation he European Bioinformatics Institute. There are no restrictions on is see by non-profit institutions as long as its content is in no wodified and this statement is not removed. Usage by and for commercintities requires a license agreement (See http://www.isb-sib.ch/announcresend an email to license@isb-sib.ch).	DEVELOPMENTAL STAGE: THE MRNA ENCODING THIS LD-NCAM IS THE MAJOR TRANSCRIPT PRESENT IN BOTH MATERNAL RNA AND IN THE EMBRYO DURING EARLY NEURAL DEVELOPMENT SIMILARITY: BELONGS TO THE IMMUNOGLOBULIN SUPERFAMILY SIMILARITY: CONTAINS 5 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS SIMILARITY: CONTAINS 2 FIBRONECTIN TYPE III-LIKE DOMAINS.	Nucleic Acids Res. 17:10321-10335(1989).	NCBI_TaxID-8355; [1] SEQUENCE FROM N.A. SEQUENCE FROM N.A. MEDLINE-90098871; PubMed-2481269; Krieg P.A., Sakaguchi D.S., Kintner C.R.; Primary structure and developmental expression of a large Cytoplasmic domain form of Xenopus laevis neural cell adhesion	an clawed frog). ;hordata; Craniata; Vertebrata; Euteleo Anura; Mesobatrachia; Pipoidea; Pipida	P16170; 01-APR-1990 (Rel. 14, Created) 01-APR-1990 (Rel. 14, Last sequence update) 15-JUN-2002 (Rel. 41, Last annotation update) Neural cell adhesion molecule 1, 180 kDa isoform precursor (N-CAM	LXENLA NCAL_XENLA STANDARD; PRT; 1088 AA.

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PVRIJLI 2
PVRI PVRI
AC Q9GI
DT 16-C
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DE POLI
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OS Sus
OC Euka
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Best Local S
Matches 89
      PVR1_PIG STANDARD; PRT; 996L76; Q96L76; 16-OCT-2001 (Rel. 40, Created) 16-OCT-2001 (Rel. 40, Last sequence upon 15-JUN-2002 (Rel. 41, Last annotation poliovirus receptor related protein 1 mediator C) (HveC) (Nectin 1).

PVRL1 OR PRR1 OR HVBC.

Sus Strofa (Pig).
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                                                                                                                                                                                                  TDAGEYFCIASNP-IGVDMQAM-YFEVQYAPKIR----GPVVVYTWEGNPVNITCEVFAH
                                                                                                                                                                                                                        DDGVPVICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDALELTCEAIGK
                                                                                                                                                                                                                                        EITLTCEA-SGDPIPSITWRTAVRNISSEATTLDGHIVVKEHIRM----SALTLKDIQY
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                                                                                                                              EFILVQADTPSS----PAIRKVEPYSSTVMIVFDEPDSTGG
                                                                                                                                               DYMLYVYDPPTTIPPPTTTTTTTTTTTTTLTIITDSRAG
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d AA; 1
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Chordata;
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26.2%;
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ed protein 1
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IG-LIKE CZ-TYPE DOMAIN 2.
IG-LIKE CZ-TYPE DOMAIN 3.
IG-LIKE CZ-TYPE DOMAIN 3.
IG-LIKE CZ-TYPE DOMAIN 4.
IG-LIKE CZ-TYPE DOMAIN 5.
FIBRONECTIN TYPE-III 1.
FIBRONECTIN TYPE-III 2.
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2; Mismatches
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Craniata;
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No. 3
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on update)
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Vertebrata;
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Euteleostomi;
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Best Local Similarity
Matches 80; Conser
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Pfam; PF00047; Ig; 2.
SMART; SM00409; IG; 3.
SMART; SM00410; IG_11ke; 1
SMART; SM00408; IGC2; 2.
SMART; SM00406; IGC; 1.
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CARBOHYD
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HSSP;
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is a functional alphaherpesvirus
virology 281:315-328(2001).
-!- FUNCTION: PROBABLY INVOLVED I
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Cohen G.H.;
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 74
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                                                 adhesion; Immunoglobulin
   TNVSISDEGRYFCQLYTDPP-
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; IPR003006;
; IPR003598;
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1176378; PubMed=11277703;
P.. Connolly S.A., Krumm
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Ig_like.
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E C2-TYPE DOMAIN 1.
E C2-TYPE DOMAIN 2.
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ES VIRUS)
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Post-processing: Minimum Match 0%
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Perfect score:
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          Published_Applications_AA:*

1: /cgn2_6/ptodata/1/pubpaa/Pi
2: /cgn2_6/ptodata/1/pubpaa/Pi
3: /cgn2_6/ptodata/1/pubpaa/Pi
3: /cgn2_6/ptodata/1/pubpaa/Ui
4: /cgn2_6/ptodata/1/pubpaa/Ui
6: /cgn2_6/ptodata/1/pubpaa/Ui
6: /cgn2_6/ptodata/1/pubpaa/Ui
6: /cgn2_6/ptodata/1/pubpaa/Ui
9: /cgn2_6/ptodata/1/pubpaa/Ui
10: /cgn2_6/ptodata/1/pubpaa/Ui
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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Sequence 61, Appl Sequence 61, Appl Sequence 61, Appl Sequence 34, Appl Sequence 20, Appl Sequence 2, Appl Sequence 2, Appl	661, 861, 886, 861, 886, 861, 886, 861, 886, 861, 886, 861, 886, 886	Description

## ALIGNMENTS

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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P2548P1C1
CURRENT APPLICATION NUMBER: US/09/944,413
CURRENT APPLICATION NUMBER: US/09/944,413
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 00/9-86
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/67,411
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,335
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,278
PRIOR APPLICATION NUMBER: 60/069,278
PRIOR APPLICATION NUMBER: 60/069,425
PRIOR APPLICATION NUMBER: 60/069,425
PRIOR FILING DATE: December 12, 1997
PRIOR APPLICATION NUMBER: 60/069,425
PRIOR FILING DATE: December 15, 1997
PRIOR APPLICATION NUMBER: 60/069,694
PRIOR FILING DATE: December 16, 1997
PRIOR APPLICATION NUMBER: 60/069,694
PRIOR FILING DATE: December 16, 1997
PRIOR APPLICATION NUMBER: 60/069,694
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Tumas,Daniel
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Godowski, Paul
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DATE:
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R APPLICATION NUMBER: (R FILING DATE: December R APPLICATION NUMBER: (R FILING DATE: December)

December

NUMBER: 60/069,702 : December 16, 1997

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OR APPLICATION NUMBER: 09/218 517
OR FILING DATE: December 22, 1998
OR APPLICATION NUMBER: 09/254,311
OR FILING DATE: March 3, 1999
OR APPLICATION NUMBER: PCT/US99/12252
OR FILING DATE: June 22, 1999
OR APPLICATION NUMBER: PCT/US99/21090
OR APPLICATION NUMBER: PCT/US99/21090
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R APPLICATION NUMBER: 60/112,850
DR FILING DATE: December 16, 1998
DR FILING DATE: December 22, 1998
DR FILING DATE: December 22, 1998
DR APPLICATION NUMBER: 60/113,296
DR FILING DATE: December 29, 1999
DR FILING DATE: July 28, 1999
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FILING DATE: May 22, 2000
APPLICATION NUMBER: PCT/US00/20710
FILING DATE: July 28, 2000
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FILING DATE: March 30,
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APPLICATION NUMBER: PCT/US00/05841
FILING DATE: March 2, 2000
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APPLICATION NUMBER: PCT/US99/30095
FILING DATE: December 16, 1999
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FILING DATE: No. US20020156004Alember
APPLICATION NUMBER: PCT/US99/28301
FILING DATE: December1, 1999
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FILING DATE: September 16, 1998
APPLICATION NUMBER: PCT/US98/25108
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APPLICATION NUMBER: 60/070,440
FILING DATE: January 5, 1998
APPLICATION NUMBER: 60/074,086
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FILING DATE:
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NUMBER: PCT/US00/04414
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NUMBER: PCT/US99/28409
NO. US20020156004Alember
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NUMBER: PCT/US01/06520
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RESULT 2
US-09-944-403-61
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PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/06935
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,278
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,425
PRIOR FILING DATE: December 12, 1997
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PRIOR APPLICATION NUMBER: 60/069,696
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PRIOR FILING DATE: December 16, 1997
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/94.
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 09/866,028
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
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                                                       APPLICATION NUMBER: 60/069,702
FILING DATE: December 16, 1970
APPLICATION NUMBER: 60/069,870
FILING DATE: December 17, 1997
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Tumas, Daniel
Wood, William
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NUMBER: 60/069,873
: December 17, 1997
NUMBER: 60/068,017
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R FILING DATE: December 16, 1998
DR APPLICATION NUMBER: 09/218,517
DR FILING DATE: December 22, 1998
DR APPLICATION NUMBER: 09/254,311
DR FILING DATE: March 3, 1999
R FILING DATE: March 3, 1999
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OF FILING DATE: February 9, 1998
OR APPLICATION NUMBER: 60/075,945
OR FILING DATE: February 25, 1998
OR APPLICATION NUMBER: 60/112,850
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 60/113,296
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OR APPLICATION NUMBER: 60/146,222
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APPLICATION NUMBER: PCT/US00/
TOTAL DATE: February 11, 200
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APPLICATION NUMBER: PCT/US00/08439
FILING DATE: MARCh 30, 2000
APPLICATION NUMBER: PCT/US00/14042
FILING DATE: May 22, 2000
APPLICATION NUMBER: PCT/US00/20710
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APPLICATION NUMBER: PCT/US99/28409
FILING DATE: No. US20020165143A1ember 30, 1999
APPLICATION NUMBER: PCT/US99/28313
FILING DATE: No. US20020165143A1ember 30, 1999
APPLICATION NUMBER: PCT/US99/28301
FILING DATE: December1, 1999
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APPLICATION NUMBER: PCT/US00/32678
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APPLICATION NUMBER: PCT/US99/30095
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APPLICATION NUMBER: PCT/US98/19330
FILING DATE: September 16, 1998
APPLICATION NUMBER: PCT/US98/25108
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P2548P1C1

CURRENT APPLICATION NUMBER: 02/09/944,896

CURRENT FILING DATE: 2001-08-31

PRIOR APPLICATION NUMBER: 60/069,334

PRIOR FILING DATE: 2001-05-25

PRIOR FILING DATE: 2001-05-25

PRIOR APPLICATION NUMBER: 60/069,334

PRIOR FILING DATE: December 11, 1997

PRIOR APPLICATION NUMBER: 60/069,278

PRIOR FILING DATE: December 11, 1997

PRIOR APPLICATION NUMBER: 60/069,278

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PRIOR APPLICATION NUMBER: 60/069,425

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PRIOR APPLICATION NUMBER: 60/069,696

PRIOR FILING DATE: December 16, 1997

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OR FILING DATE: December 22, 1998
OR APPLICATION NUMBER: 09/254,311
OR FILING DATE: March 3, 1999
OR APPLICATION NUMBER: PCT/US99/1225
OR FILING DATE: June 22, 1999
OR APPLICATION NUMBER: PCT/US99/21090
OR APPLICATION NUMBER: PCT/US99/21090
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APPLICATION NUMBER: PCT/US00/05841
FILING DATE: March 2, 2000
APPLICATION NUMBER: PCT/US00/08439
FILING DATE: March 30, 2000
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APPLICATION NUMBER: PCT/US01/06520
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FILING DATE: December1, 199
APPLICATION NUMBER: PCT/US99/30095
FILING DATE: December 16, 1999
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APPLICATION NUMBER: PCT/US99/28409
FILING DATE: NO. US20020168715A1ember
APPLICATION NUMBER: PCT/US99/28313
FILING DATE: NO. US20020168715A1ember
APPLICATION NUMBER: PCT/US99/28301
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APPLICATION NUMBER: PCT/US98,
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                    TWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIP
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   TWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIP
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: December 22, 1998
: NUMBER: 60/146,222
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US-09-866-028-61
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NUMBER OF SEQ ID NOS: 120
SEQ ID NO 61
LENGTH: 440
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APPLICANT:
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CURRENT FILING DATE: 2001-05-25
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mes 336; Conserv
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 PPTTTTTTTTTTTLTIITDSRAGEEGSIRAVDH
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                                                                     TWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIP
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Godowski, Paul
Grimaldi, Christopher
Gurney, Austin
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Tumas,Daniel
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Kljavin, Ivar
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Pred. No. 1.7e-105;
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RESULT 5
US-09-944-449-61
US-09-944-449-61
; Sequence 61, Application US/09944449
; Patent No. US20020102647A1

GENERAL INFORMATION:

APPLICANT:

Baker, Kevin Botstein, David

Eaton, Dan

APPLICANT:

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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 09/866,07
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,47
PRIOR FILING DATE: December 3, 1997
PRIOR PRIOR PRIOR NUMBER: 60/069,37
PRIOR FILING DATE: December 11, 1997
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CURRENT APPLICATION NUMBER: US/09/944,449
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES
TITLE OF INVENTION: ACIDS ENCODING THE SAME
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OR APPLICATION NUMBER: PCT/US98/1930
OR FILING DATE: September 16, 1998
OR APPLICATION NUMBER: PCT/US98/25108
OR FILING DATE: December 1, 1998
OR APPLICATION NUMBER: 09/216, 021
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 09/218, 517
OR FILING DATE: December 22, 1998
OR APPLICATION NUMBER: 09/24, 311
OR FILING DATE: March 3, 1999
OR APPLICATION NUMBER: PCT/US99/1252
OR APPLICATION NUMBER: PCT/US99/1252
OR APPLICATION NUMBER: PCT/US99/1209
OR APPLICATION NUMBER: PCT/US99/21090
OR FILING DATE: June 22, 1999
OR APPLICATION NUMBER: PCT/US99/21090
OR FILING DATE: September 15, 1999
                                                                                                                                                                                                                                                                                OR APPLICATION NUMBER: 60/075,947
OR APPLICATION NUMBER: 60/069,694
OR FILING DATE: December 16, 1997
OR APPLICATION NUMBER: 60/069,694
OR FILING DATE: December 16, 1997
OR APPLICATION NUMBER: 60/069,870
OR APPLICATION NUMBER: 60/069,870
OR APPLICATION NUMBER: 60/069,873
OR FILING DATE: December 17, 1997
OR APPLICATION NUMBER: 60/070,440
OR FILING DATE: December 18, 1997
OR APPLICATION NUMBER: 60/070,440
OR FILING DATE: FEBRUARY 9, 1998
OR APPLICATION NUMBER: 60/074,092
OR FILING DATE: FEBRUARY 9, 1998
OR APPLICATION NUMBER: 60/075,945
OR APPLICATION NUMBER: 60/113,985
OR APPLICATION NUMBER: 60/113,1996
OR APPLICATION NUMBER: 60/146,222
OR FILING DATE: December 2, 1998
OR APPLICATION NUMBER: 60/146,222
OR FILING DATE: December 22, 1998
OR APPLICATION NUMBER: 60/146,222
OR FILING DATE: December 20, 1998
OR APPLICATION NUMBER: 60/146,222
OR FILING DATE: December 30/146,222
OR FILING DATE: DECEMBER: 60/146,222
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FILING DATE: December 11, 1997
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Tumas, Daniel
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E: December 12, 1997
N NUMBER: 60/069,696
E: December 16, 1997
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Best Local Similarity
Matches 336; Conserv
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PRIOR FILING DATE: N
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IOR FILING DATE: February 11, 2000
IOR APPLICATION NUMBER: PCT/US00/04414
IOR APPLICATION NUMBER: PCT/US00/04414
IOR FILING DATE: February 22, 2000
IOR APPLICATION NUMBER: PCT/US00/05841
IOR FILING DATE: March 2, 2000
IOR APPLICATION NUMBER: PCT/US00/14042
IOR FILING DATE: March 30, 2000
IOR FILING DATE: March 30, 2000
IOR FILING DATE: March 30, 2000
IOR FILING DATE: May 22, 2000
IOR APPLICATION NUMBER: PCT/US00/20710
IOR APPLICATION NUMBER: PCT/US00/20710
IOR APPLICATION NUMBER: PCT/US00/23678
IOR APPLICATION NUMBER: PCT/US00/32678
IOR APPLICATION NUMBER: PCT/US00/32678
IOR APPLICATION NUMBER: PCT/US00/32678
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No. US20020102647Alember
NUMBER: PCT/US99/28313
No. US20020102647Alember
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Sequence 61, Application US/09944457

Patent No. US20020110899A1

GENERAL INFORMATION:
APPLICANT: Baker, Kevin
APPLICANT: Botstein, David
APPLICANT: Eaton, Dan

APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Gerritsen, Mary
Gerritsen, Mary
Goddard, Audrey
Godowski, Paul
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APPLICANT:
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OR APPLICATION NUMBER: 60/067,411
OR FILING DATE: December 3, 1997
OR APPLICATION NUMBER: 60/069,335
OR FILING DATE: December 11, 1997
OR APPLICATION NUMBER: 60/069,278
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OR FILING DATE: December 12, 1997
OR APPLICATION NUMBER: 60/069,694
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FILING DATE: February 9, 1998
FILING DATE: February 0
                                    FILING DATE: September 15, 1999
APPLICATION NUMBER: PCT/US99/28409
FILING DATE: NO. US200/20110859Alember
APPLICATION NUMBER: PCT/US99/28313
FILING DATE: No. US200/20110859Alember
APPLICATION NUMBER: PCT/US99/28301
FILING DATE: December1, 1999
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FILING DATE: March 3, 1999
APPLICATION NUMBER: PCT/US99/12252
FILING DATE: June 22, 1999
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APPLICATION NUMBER: 09/218,517
FILING DATE: December 22, 1998
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APPLICATION NUMBER: 60/146,222
FILING DATE: July 28, 1999
APPLICATION NUMBER: PCT/US98/19330
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APPLICATION NUMBER: 60/113,296
FILING DATE: December 22, 1998
APPLICATION NUMBER: 60/146,222
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APPLICATION NUMBER: 60/070,440
FILING DATE: January 5, 1998
APPLICATION NUMBER: 60/074,086
FILING DATE: February 9, 1998
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APPLICATION NUMBER: 60,069,702
FILING DATE: December 16, 1997
APPLICATION NUMBER: 60,069,870
FILING DATE: December 17, 1997
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 APPLICATION FILING DATE
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APPLICATION NUMBER: 09/216,021
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Hillan,Kenneth
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Napier, Mary
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                   NUMBER: PCT/US99/30095
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Gerritsen, Mary
Goddard, Audrey
Godowski, Paul
Grimaldi, Christopher
Gurney, Austin
Hillan, Kenneth
Kljavin, Ivar

Ferrara, Napoleone Filvaroff, Ellen

Eaton, Dan Botstein, David

Kevin

Napier, Mary Roy, Margaret Tumas, Daniel

Wood, William

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; ORGANISM: HOMO US-09-944-457-61
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SEQ ID N
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TYPE: PRT
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OR FILING DATE: MAICH 30, 2000
OR APPLICATION NUMBER: PCT/US00/14042
OR FILING DATE: MAY 22, 2000
OR APPLICATION NUMBER: PCT/US00/20710
OR FILING DATE: July 28, 2000
OR FILING DATE: July 28, 2000
OR APPLICATION NUMBER: PCT/US00/32678
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Local Similarity
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FILING DATE: February 22, 2000
APPLICATION NUMBER: PCT/US00/05841
FILING DATE: March 2, 2000
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                                                                TWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIP
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PRIOR APPLICATION NUMBER: 60/069
PRIOR FILING DATE: December 11, 1
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                                OR APPLICATION NUMBER: PCT/US00/03565
OR APPLICATION NUMBER: PCT/US00/03565
OR FILING DATE: February 11, 2000
OR FILING DATE: February 22, 2000
IOR APPLICATION NUMBER: PCT/US00/05841
IOR APPLICATION NUMBER: PCT/US00/05841
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OR APPLICATION NUMBER: 60/070,440
OR FILING DATE: January 5, 1998
OR APPLICATION NUMBER: 60/074,086
OR FILING DATE: February 9, 1998
OR APPLICATION NUMBER: 60/074,092
OR FILING DATE: February 9, 1998
OR APPLICATION NUMBER: 60/075,945
OR FILING DATE: February 25,1998
OR APPLICATION NUMBER: 60/112,850
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 60/113,296
OR FILING DATE: December 22, 1998
OR APPLICATION NUMBER: 60/113,296
OR APPLICATION NUMBER: 60/113,296
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OR APPLICATION NUMBER: 60/146,222
OR FILING DATE: JULY 28, 1999
OR APPLICATION NUMBER: PCT/US98/19330
OR FILING DATE: September 16, 1998
OR APPLICATION NUMBER: PCT/US98/25108
OR APPLICATION NUMBER: PCT/US98/25108
OR APPLICATION NUMBER: 09/216,021
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DR FILING DATE: December 16, 1997
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FILING DATE:
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APPLICATION NUMBER: 09/218,517
FILLING DATE: December 22, 1998
APPLICATION NUMBER: 09/254,311
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ETILING DATE: June 22, 1999
APPLICATION NUMBER: PCT/US99/21090
FILING DATE: September 15, 1999
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FILING DATE: December 11, 199
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APPLICATION NUMBER: PCT/US99/30095
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FILING DATE: No. US20020115145Alember
APPLICATION NUMBER: PCT/US99/28301
FILING DATE: December1, 1999
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APPLICATION NUMBER: PCT/US99/28409
FILING DATE: No. US20020115145Alember
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NUMBER: 60/069,425
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PRIOR TILING DATE: MBy 22, 2000
PRIOR APPLICATION NUMBER: PCT/US00/20710
PRIOR FILING DATE: July 28, 2000
PRIOR APPLICATION NUMBER: PCT/US00/32678
PRIOR PILING DATE: December 1, 2000
PRIOR APPLICATION NUMBER: PCT/US01/06520
CURRENT APPLICATION NUMBER: US/09/945,587 CURRENT FILING DATE: 2001-09-26 PRIOR APPLICATION NUMBER: 09/866,028 PRIOR FILING DATE: 2001-05-25 PRIOR APPLICATION NUMBER: 60/067,411
                                                                                                         TITLE OF INVENTION: SECRETED TITLE OF INVENTION: ACIDS ENTITLE REFERENCE: P2548P1C1
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Tumas, Daniel
Wood, William
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Gerritsen, Mary
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Grimaldi,Christopher
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OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 09/218,517
OR FILING DATE: December 22, 1998
OR APPLICATION NUMBER: 09/254,311
OR FILING DATE: March 3, 1999
OR APPLICATION NUMBER: PCT/US99/1252
OR APPLICATION NUMBER: PCT/US99/1252
OR APPLICATION NUMBER: PCT/US99/21090
OR APPLICATION NUMBER: PCT/US99/21090
OR APPLICATION NUMBER: PCT/US99/21090
OR APPLICATION SEPTEMBER: 15, 1999
OR FILING DATE: September 15, 1999
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R FILING DATE: December 16, 1998
DR APPLICATION NUMBER: 60/113,296
DR FILING DATE: December 22, 1998
DR APPLICATION NUMBER: 60/146,222
DR FILING DATE: July 28, 1999
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APPLICATION NUMBER: 60/069,870
                                                                                                                                                                   FILING DATE:
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                                                     N NUMBER: PCT/US00/14042
E: May 22, 2000
N NUMBER: PCT/US00/20710
N NUMBER: PCT/US00/20710
PE: July 28, 2000
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N NUMBER: PCT/US00/05841

N NUMBER: PCT/US00/08439

N NUMBER: PCT/US00/08439

E: March 30, 2000
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: NO. US20020127643A1ember
NUMBER: PCT/US99/28313
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:: September 16, 1998
NUMBER: PCT/US98/25108
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APPLICANT: Tunas, Daniel
APPLICANT: Wood, William
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND
TITLE OF INVENTION: SECRETED AND THE SAME
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P2548P1C1
CURRENT APPLICATION NUMBER: U5/09/945,015
CURRENT APPLICATION NUMBER: 09/866,028
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069335
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,278
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PRIOR APPLICATION NUMBER: 60/069,278
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; TYPE: PRT
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US-09-945-587-61
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Grimaldi, Christopher
Gurney, Austin
Hillan, Kenneth
Kljavin, Ivar
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Tumas,Daniel
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; TYPE: PRT; ORGANISM: HOMO US-09-945-015-61
                                                                                                                                                      NUMBER OF S
SEQ ID NO 61
  Query Match
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**KIOR APPLICATION NUMBER: PCT/USO1,
PRIOR FILING DATE: February 28, 200
NUMBER OF SEQ ID NOS: 100
EQ ID NO 61
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OR APPLICATION NUMBER: 60/074,086
OR FILING DATE: February 9, 1998
OR APPLICATION NUMBER: 60/074,092
OR FILING DATE: February 9, 1998
OR APPLICATION NUMBER: 60/075,945
OR FILING DATE: February 25, 1998
OR APPLICATION NUMBER: 60/112,850
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 60/113,296
OR FILING DATE: December 22, 1998
OR APPLICATION NUMBER: 60/146,222
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: PCT/US98/19330
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: PCT/US98/19330
OR FILING DATE: December 1, 1998
OR APPLICATION NUMBER: 09/216,021
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 09/218,517
OR FILING DATE: December 22, 1998
OR APPLICATION NUMBER: 09/218,517
OR FILING DATE: December 22, 1998
OR APPLICATION NUMBER: 09/218,517
OR FILING DATE: December 22, 1998
OR APPLICATION NUMBER: 09/218,517
OR FILING DATE: March 3, 1999
OR APPLICATION NUMBER: PCT/US99/12252
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FILING DATE: September 15, 1999
APPLICATION NUMBER: PCT/US99/28409
FILING DATE: NO. US20020132768Alember 30, 1999
APPLICATION NUMBER: PCT/US99/28313
FILING DATE: NO. US20020132768Alember 30, 1999
APPLICATION NUMBER: PCT/US99/28301
FILING DATE: NO. US20020132768Alember 30, 1999
APPLICATION NUMBER: PCT/US99/28301
FILING DATE: December1, 1999
APPLICATION NUMBER: PCT/US99/28301
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APPLICATION NUMBER: 60/069,873
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FILING DATE: May 22, 2000
APPLICATION NUMBER: PCT/US00/20710
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APPLICATION NUMBER: 60/069,702
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APPLICATION NUMBER: PCT/US00/04414
FILING DATE: February 22, 2000
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Length 440;
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APPLICANT: WOOD, WILLIAM
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P2548P1C1
CURRENT APPLICATION NUMBER: US/09/944,396
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 09/866,028
PRIOR APPLICATION NUMBER: 09/867,411
PRIOR FILING DATE: December 3, 1997
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR APPLICATION NUMBER: 60/069335
PRIOR APPLICATION NUMBER: 60/069335
PRIOR FILING DATE: December 11, 1997
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                                                                                                                       APPLICATION NUMBER: 60/069,278 FILING DATE: December 11, 1997 APPLICATION NUMBER: 60/069,425
APPLICATION FILING DATE
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Godowski, Paul
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Filvaroff, Ellen
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                                                                   NUMBER: 60/069,425
December 12, 1997
NUMBER: 60/069,696
December 16, 1997
                  NUMBER:
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LENGTH: 44
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NUMBER OF SEQ ID NO
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APPLICATION NUMBER: PCT/US00/20710
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FILING DATE: September 15, 1999
APPLICATION NUMBER: PCT/US99/28409
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FILING DATE: March 3, 199
APPLICATION NUMBER: PCT/US99/12252
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APPLICATION NUMBER: PCT/US98/25108
FILING DATE: December 1, 1998
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APPLICATION NUMBER: 60/075,
FILING DATE: February 25, 1
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                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: PCT/US00/32678
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APPLICATION NUMBER: 09/218,517
FILING DATE: December 22, 1998
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APPLICATION NUMBER: PCT/US98/19330
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                                     IPTGDGQNLFTKDVTVIEGEVATISCQVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQ
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NUMBER: 09/216,021
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CURRENT FILLING DATE: 2001-08-31
PRIOR APPLICATION NUMBER: 09/866,028
PRIOR FILING DATE: 2001-05-25
PRIOR PPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069335
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DR APPLICATION NUMBER: 60/069,696
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APPLICATION NUMBER: 60/069,873
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Tumas, Daniel
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OR FILING DATE: MARCH 30, 2000
OR APPLICATION NUMBER: PCT/US00/14042
IOR FILING DATE: MAY 22, 2000
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OR APPLICATION NUMBER: 09/218,517
OR FILING DATE: December 22, 1998
OR APPLICATION NUMBER: 09/254,311
OR FILING DATE: MATCh 3, 1999
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APPLICATION NUMBER: PCT/US00/04414
FILING DATE: February 22, 2000
APPLICATION NUMBER: PCT/US00/05841
FILING DATE: March 2, 2000
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APPLICATION NUMBER: PCT/US99/30095
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APPLICATION NUMBER: PCT/US00/03565
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Mismatches 0;
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PRIOR FILING DATE: February 9, 199
PRIOR APPLICATION NUMBER: 60/075, 9
PRIOR FILING DATE: February 25, 199
PRIOR APPLICATION NUMBER: 60/112, 89
PRIOR FILING DATE: December 16, 199
PRIOR APPLICATION NUMBER: 60/113, 99
PRIOR FILING DATE: December 22, 199
PRIOR APPLICATION NUMBER: 60/146, 22
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PRIOR FILLING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,425
PRIOR FILING DATE: December 12, 1997
PRIOR APPLICATION NUMBER: 60/069,696
PRIOR FILING DATE: December 16, 1997
PRIOR FILING DATE: December 16, 1997
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 09/866,028
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
PRIOR FILING DATE: DECEMBER: 60/069,334
PRIOR APPLICATION NUMBER: 60/069,334
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PRIOR APPLICATION UNMER: 60/069,873
PRIOR FILING DATE: December 17, 1997
PRIOR APPLICATION NUMBER: 60/068,017
PRIOR APPLICATION NUMBER: 60/068,017
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R FILING DATE: February 9, 1,7

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PAR APPLICATION UNMBER: 60/074,092

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PAR APPLICATION UNMBER: 60/075,945
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OR FILLING DATE: December 16, 1997
OR APPLICATION NUMBER: 60/069,702
OR FILLING DATE: December 16, 1997
OR APPLICATION NUMBER: 60/069,870
OR FILING DATE: December 17, 1997
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IOR FILING DATE: NO. US20020142419Alember 30
IOR APPLICATION NUMBER: PCT/US99/28313
IOR FILING DATE: NO. US20020142419Alember 30
IOR APPLICATION NUMBER: PCT/US99/28301
IOR FILING DATE: December1, 1999
IOR FILING DATE: December 16, 1999
IOR FILING DATE: December 16, 1999
IOR APPLICATION NUMBER: PCT/US99/30095
IOR APPLICATION NUMBER: PCT/US90/03565
IOR APPLICATION NUMBER: PCT/US90/04414
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IOR FILING DATE: February 22, 2000
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OR FILING DATE: MAICH 2, 2000
OR APPLICATION NUMBER: PCT/US00/08439
OR FILING DATE: MAICH 30, 2000
OR APPLICATION NUMBER: PCT/US00/14042
OR FILING DATE: May 22, 2000
OR APPLICATION NUMBER: PCT/US00/20710
OR APPLICATION NUMBER: PCT/US00/20710
OR FILING DATE: July 28, 2000
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DR FILING DATE: December 22, 1998
DR APPLICATION NUMBER: 09/254,311
DR FILING DATE: March 3, 1929
DR APPLICATION NUMBER: PCT/US99/12252
DR FILING DATE: June 22, 1999
DR APPLICATION NUMBER: PCT/US99/21090
DR APPLICATION NUMBER: PCT/US99/21090
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RESULT 13
US-09-943-762-61
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PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069335
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069278
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PRIOR FILING DATE: December 11, 1997
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APPLICANT: Roy, Margaret
APPLICANT: Tumas, Daniel
APPLICANT: Wood, William
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P2548P1C1
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                                    A APPLICATION NUMBER: 6U/12 PT TILING DATE: JULIY 28, 1999
TO STLING DATE: JULIY 28, 1999
TO NUMBER: PCT/US98/19330
TO NUMBER: PCT/US98/19330
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TO 25108
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OR FILING DATE: December 17, 1997
OR APPLICATION NUMBER: 60/069,873
OR TILING DATE: December 17, 1997
OR APPLICATION NUMBER: 60/068,017
OR APPLICATION NUMBER: 60/068,017
OR FILING DATE: December 18, 1997
OR APPLICATION NUMBER: 60/070,440
OR FILING DATE: January 5, 1998
OR FILING DATE: January 5, 1998
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APPLICATION NUMBER: 60/069,702
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APPLICATION NUMBER: 09/216,
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WHIMBER: 60/070,440

TE: January 5, 1998

WHIMBER: 60/074,086

TE: February 9, 1998

WHIMBER: 60/074,092

TE: February 9, 1998

TE: February 25, 1998

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N NUMBER: 60/113,296
E: December 22, 1998
N NUMBER: 60/146,222
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NUMBER: 60/069,694
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RESULT 14
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; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin
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OR APPLICATION NUMBER: PCT/US00/03565
OR APPLICATION NUMBER: PCT/US00/04414
OR APPLICATION NUMBER: PCT/US00/04414
OR FILING DATE: February 22, 2000
OR APPLICATION NUMBER: PCT/US00/05841
OR FILING DATE: March 2, 2000
OR APPLICATION NUMBER: PCT/US00/08439
OR FILING DATE: March 30, 2000
OR FILING DATE: March 30, 2000
OR APPLICATION NUMBER: PCT/US00/14042
OR FILING DATE: May 22, 2000
OR APPLICATION NUMBER: PCT/US00/14042
OR FILING DATE: May 22, 2000
OR APPLICATION NUMBER: PCT/US00/20710
OR FILING DATE: May 22, 2000
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FILING DATE: February 28
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FILING DATE: No. US20020142958Alember 30, 1999
APPLICATION NUMBER: PCT/US99/28313
FILING DATE: No. US20020142958Alember 30, 1999
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NUMBER: 09/254,311
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PRIOR FILLING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILLING DATE: December 3, 1997
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILLING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILLING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,425
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PRIOR APPLICATION NUMBER: 60/069,425
PRIOR FILLING DATE: December 16, 1997
PRIOR APPLICATION NUMBER: 60/069,694
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PRIOR APPLICATION NUMBER: 60/069,702
PRIOR FILLING DATE: December 17, 1997
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PRIOR APPLICATION NUMBER: 60/069,870
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PRIOR APPLICATION NUMBER: 60/069,873
PRIOR FILLING DATE: December 17, 1997
PRIOR APPLICATION NUMBER: 60/069,873
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PRIOR APPLICATION NUMBER: 60/068,017
PRIOR FILLING DATE: December 18, 1997
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Godowski, Paul
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                                                            E: February 9, 1998

N NUMBER: 60/074,092

TE: February 9, 1998

N NUMBER: 60/075,945

N NUMBER: 60/112,850

N NUMBER: 60/113,296

TE: December 16, 1998

TE: NUMBER: 60/113,296

TE: December 22, 1998

N NUMBER: 60/146,222

TE: July 28, 1999
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E: December 16, 1998
N NUMBER: 09/218,517
E: December 22, 1998
N NUMBER: 09/254,311
                               NUMBER: PCT/US99/12252
: June 22, 1999
NUMBER: PCT/US99/21090
: September 15, 1999
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                               APPLICANT: Baker, MAPPLICANT: Botstein APPLICANT: Beton, I APPLICANT: Ferrara APPLICANT: Filvarra APPLICANT: Goddard APPLICANT: Goddard
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No. US20020150976A1
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APPLICATION NUMBER: PCT/US00/14042
FILING DATE: May 22, 2000
APPLICATION NUMBER: PCT/US00/20710
FILING DATE: July 28, 2000
APPLICATION NUMBER: PCT/US00/32678
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APPLICATION NUMBER: PCT/US00/04414
FILING DATE: February 22, 2000
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APPLICATION NUMBER: PCT/US00/03565
FILING DATE: February 11, 2000
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Goddard, Audrey
 Godowski, Paul
Grimaldi, Christopher
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PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069335
PRIOR APPLICATION NUMBER: 60/0693278
PRIOR APPLICATION NUMBER: 60/069,278
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APPLICANT: Wood, William
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
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OR APPLICATION NUMBER: 60,074,092
OR APPLICATION NUMBER: 60,075,945
OR APPLICATION NUMBER: 60,075,945
OR FILING DATE: Pebruary 25, 1998
OR APPLICATION NUMBER: 60,7112,850
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 60,7113,296
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APPLICATION NUMBER: 60/146,222
FILING DATE: July 28, 1999
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APPLICATION NUMBER: 60/074,08
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PRIOR FILING DATE: March 2, 2000
PRIOR APPLICATION NUMBER: PCT/US00/08439
PRIOR FILING DATE: March 30, 2000
PRIOR APPLICATION NUMBER: PCT/US00/14042
PRIOR FILING DATE: May 22, 2000
PRIOR APPLICATION NUMBER: PCT/US00/20710
PRIOR APPLICATION NUMBER: PCT/US00/20710
PRIOR FILING DATE: July 28, 2000
PRIOR FILING DATE: December 1, 2000
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PRIOR APPLICATION NUMBER: PCT/US01/06520
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Tang YT, Liu C, 2) Cao Y, Drmanac RA,

Zhou P, Qian XB, Wa 1, Zhang J, Werhman

Wang Z, nan T;

Chen , Z

Asundi V;

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RESULT 1
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25-JAN-2000; 2000US-0491404.
17-JUL-2000; 2000US-0617746.
03-AUG-2000; 2000US-0631451.
15-SEP-2000; 2000US-0663870
                                                                                                                                                                                                               Human; sheep; pig; cow; fruit fly; yeast; hamster; macaque; horse; tomato; monkey; dog; sea urchin; expressed sequence tag; EST; diagnostics; forensic test; gene mapping; genetic disorder; blodiversity; gene therapy; nutrition.
                                                                                                                                              02-AUG-2001.
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                                          (HYSE-) HYSEQ INC
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                                                                                                                                                                                                                                                                                                                                                                   The present invention describes nucleic acids encoding PRO secreted and transmembrane proteins used therapeutically. The PRO proteins have cytostatic, anti-inflammatory, anti-proliferative and immunosuppressive activity. The proteins and polynucleotides can be used in therapy, identification of homologues, raising antibodies and design of probes and primers. They can be used in a range of diseases related to proteins that they have homology with, e.g. a PRO protein having homology to complement proteins may be used in inflammatory responses.
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                                                                     Baker KP,
Gerritsen N
Hillan KJ,
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Peptide
      New human nucleic acids encoding secreted and transmembrane polypeptides, designated as PRO polypeptides, useful as pharmaceutical and diagnostic agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRO; membrane bound protein; secreted protein; PRO357; PRO327; PRO43; PRO715; PRO241; PRO323; PRO299; PRO233; PRO344; PRO347; PRO355; PRO355; PRO356; transmembrane polypeptide; antibody; screening; detection; inhibition; probe; primer; human.
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DB; AAA49563.
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J. Kljavin ?
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98US-0112850.
98US-0113296.
                                                                                                                                                                99WO-US28301
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427..433
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428..432
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402..4
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376..382
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372..393
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319..328
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65..69
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9..15
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                                                                     Eaton DL, Ferrara N, Filvaroff E;
A, Godowski PJ, Grimaldi CJ, Gurn
Napier MA, Roy MA, Tumas D, Woo
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RESULT 4
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ID AAU2
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XX PRO
DE Huma
XX PRO
CKW dog;
KW block
KW adre
XX WO2C
XX 20-5
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Best Local 9
01-MAR-2000;
02-MAR-2000;
03-MAR-2000;
06-MAR-2000;
14-MAR-2000;
14-MAR-2000;
15-MAR-2000;
21-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New human nucleic acids encoding secreted and transmembrane polypeptides which are designated as PRO polypeptides are described The membrane-bound proteins have various industrial applications, including as pharmaceutical and diagnostic agents. The membrane-bound proteins can also be employed for screening of potential peptide or small molecule inhibitors of the relevant receptor/ligand interaction Anti-PRO antibodies are useful for the affinity purification of PRO from recombinant cell culture or natural sources.
                                                                                                                                                                                                    PRO polypeptide; mammal; tumour; cancer; human; cattle; horse; sheep; dog; cat; plg; goat; rabbit; tumour necrosis factor alpha; TNF-alpha; blood; chondrocyte cell; cell proliferation; cell differentiation; colon; adrenal; lung; breast; prostate; rectum; cervix; liver; genetic disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim
                                                                                                         28-FEB-2001; 2001WO-US06520
                                                                                                                                                         WO200168848-A2
                                                                                                                                                                                                                                                               Human PRO polypeptide sequence
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2000WO-US05601.
2000WO-US05841.
2000WS-187202P.
2000US-186968P.
2000US-189320P.
2000US-189320P.
2000WG-189320P.
2000WG-18938P.
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                                                                                                                                                   Sequences AAU29024-AAU29328 represent PRO polypeptides of the invention.

The PRO polypeptides and their associated nucleic acids can be used to detect the presence of a tumour in a mammal by comparing the level of expression of a PRO polypeptide in a test sample of cells from the animal and a control sample of normal cells, whereby a higher level of expression in the test sample indicates the presence of a tumour in the mammals include dogs, cats, cattle, horses, sheep, pigs, goats and rabbits but are preferably human. The polypeptides can be used to stimulate tumour necrosis factor (TNF) alpha release from human blood, when contacted with it. A specific polypeptide can be used to stimulate the proliferation or differentiation of chondrocyte cells. The PRO proteins can be used to determine the presence of tumours and also susceptibility to tumour development, particularly adrenal, lung, colon, consisted. The oligonucleotide probes specific for the PRO nucleic acids of an objects. The oligonucleotide probes specific for the PRO nucleic acids can be used for generation and side of individuals with contact disparation.
                                                         Matches
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Best Local 9
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18 - APR - 2000

25 - APR - 2000

27 - AAY - 2000

27 - AAY - 2000

28 - JUN - 2000

29 - JUN - 2000

20 - JUN - 2000

21 - AUG - 2000

22 - AUG - 2000

23 - AUG - 2000

24 - AUG - 2000
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28 - MAR - 2000;
29 - MAR - 2000;
30 - MAR - 2000;
04 - APR - 2000;
04 - APR - 2000;
01 - APR - 2000;
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2000US-191148P
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2000US-19400P
2000US-195075P
2000US-196187P
2000US-19630P
2000US-19630P
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2000US-199550P
2000WO-US14042
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2000WO-US13705
2000WO-US23328
2000WO-US23328
2000WO-US23328
2000WO-US33678
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Watanabe CK,
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Pred. No. 2.1
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RESULT 5
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                                                                                                                         The present invention relates to 12 secreted human proteins and the nucleotide sequences encoding them. The polynucleotide sequences give in AAA80606-A80623 encode the 12 secreted protein sequences given in AAB25576-B25593. The human secreted proteins have various activities
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antirheumatic, dermatological; antiproliferative; antiarteriosclerotic; anticancer; vulnerary; antiviral; antibacterial; antifungal; immune disorder; Addison's disease; rheumatoid arthritis; dermatitis; multiple sclerosis; inflammatory disorder; inflammatory bowel disease; crohn's disease; nephritis; hyperproliferative disorder; cardiovascular disorder; coronary arteriosclerosis; myocarditis; cancer
                                                                                                                                                                                                                                                                                                                                  Isolated nucleic acid molecules encoding human secreused for the prevention, amelioration and treatment inflammatory, hyperproliferative and cardiovascular
                       antiproliferative;
                                                  anti-inflammatory;
                                                                  activities of the proteins include: immunosuppressant
                                                                                                 dependent on the tissues in which they are expressed.
                                                                                                                                                                                                                                                             Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-387742/33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HUMA-) HUMAN GENOME SCI INC
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                                                                                                                                                                                                                                                                                                                and infectious diseases
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                       antiarthritic; antirheumatic, dermatological;
antiarteriosclerotic; anticancer; vulnerary;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JJ,
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tment of autoimmune,
cular disorders, car
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RESULT 6
AAY94341
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Best Local Sim
Matches 336;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hyperproliferative disorders such as paraproteinaemias and purpura; cardiovascular disorders e.g. coronary arteriosclerosis and myocarditis; cancer e.g. melanoma and lymphoma. The proteins and polynucleotide sequences may also be used in wound healing and the treatment of infectious diseases. The human secreted protein gene #11 and protein sequences are represented in sequences AAA80616 and AAB25586. Sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polypeptides, agonists and antagonists may be used to treat prevent and/or diagnose various disease, disorders and conditions examples which include: immune disorders e.g. Addison's disease, rheumatoid arthritis, dermatitis, and multiple sclerosis; inflammatory disorder e.g. inflammatory bowel disease, Crohn's disease and nephritis;
            Domain
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                                                      Domain
                                                                                                               čey
                                                                                                                                                       neuroprotective; nootropic; anticonvulsant; cancer; leukaemia melanoma; rheumatoid arthritis; asthma; atherosclerosis; akat Alzheimer's diseases; multiple sclerosis; epilepsy.
                                                                                                                                                                                                                        Human cell surface receptor protein
                                                                                                                                                                                                                                                 22-AUG-2000
                                                                                                                                                                                                                                                                                          AAY94341 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                            Protein
                                                                                                                                   Homo sapiens
                                                                                                                                                                                                     Human; HCSRP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA80677-A80682 represent genes related to the secreted protein
                                                                                                  Pept1de
                                                                                                                                                                                        Human; HCSRP; cytostatic; antiarthritic; antirheumatic; antiasthmatic;
immunosuppressive; antiarteriosclerotic; antibacterial; antiparasitic;
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45..442
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57..126
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159..22
                                                                                                             Location/Qualifiers
                      /label-
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           315
                      Immunoglobulin_domain
                                                                 HCSRP-8
                                                                                      Signal_peptide
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Pred. No. 2.1e-122;
; Mismatches 0;
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Claim
                                                                                                                                         New human cell surface receptor protein and diagnosis, prevention and treatment of cance infection and neuronal disorders -
                                                                                                                                                                                         WPI; 2000-376546/32.
N-PSDB; AAA27051.
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07-DEC-1998;
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                                                                                                                                                                                                                        Hillman JL,
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                                                                                                                     1; Page 81-82; 97pp; English.
                                                                                                                                                                                                                                    Corley
                                                                                                                                                                                                                        rley NC, G
Bandman O,
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98US-0206647.
99US-0123404.
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O, Azimzai Y,
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The present sequence is a novel human cell surface receptor protein (HCSRP) designated HCSRP-8. The nucleotide sequence was identified in Incyte Clone 312256 from the cDNA library LUNGNOTO2, which was made from RNA isolated from lung tissue. A number of Incyte Clones were used to assemble the consensus sequence. BLAST analysis showed that the sequence is homologous to immuno-superfamily protein B1 g3779242. HCSRP and its antagonist are useful for preventing or treating disorders associated with decreased or increased expression or activity of HCSRP. Such disorders include cancers such as leukaemia and melanoma, immune
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, Au-Young J;
                                                                                                                                                                                                                                                                                                                                                                                    n and polynucleotide useful cancer, immune disorders,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    site"
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RESULT 7
AAY45092
ID AAY4
XX AAY4
AC AAY4
XX Huma
XX Huma
XX Lymp
KW Lymp
KW B7-LL
KW inmu
KW Orga
XX Homc
XX Homc
FT Poons
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    bacterial and parasitic infections and neuronal disorders such as akathesia, Alzheimer's disease, multiple sclerosis and epilepsy. Polynucleotides encoding HSCRPs may be used as hybridisation probes diagnose these conditions. Anti-HCSRP antibodies may be used as antagonists, as a targeting or delivery mechanism for bringing pharmaceutical agents into contact with cells or tissues expressing HCSRP and for diagnosis of HCSRP-related disorders. HCSRP and its catalytic or immunogenic fragments are useful for drug screening usilibraries of compounds.
                                                                                                                                                                                                                biological activity; quality control reagent; treatme
immune system disorder; autoimmune; viral infection;
organ transplant rejection; bone marrow; modulator;
                                                                                                                                                                                                                                                      Lymphoid derived dendritic cell adhesion molecule; LDCAM; human; B7-L1; T cell proliferation; natural killer cell; NK; tumour cel
                          Modified-site
                                                   Modified-site
                                                                             Modified-site
                                                                                                                             Peptide
                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                Human
                                                                                                                                                                                                                                                                                                                        31-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                         AAY45092 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                AAY45092;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GEEIEVNCTAMASKPATTIRWFKGNTELKGKSEVEEWSDMYTVTSQLMLKVHKEDDGVPV
                                                                                                                                                                                                                                                                                                                                                                                                                                                  PPTTTTTTTTTTTILTIITDSRAGEEGSIRAVDH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LLNFSSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVE
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                                                                                                                                                                                                                                                                                               lymphoid
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                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                            derived dendritic cell adhesion molecule.
                         /note= "N
113..115
                                                  /note= "J
 /note=
165..1
                                                                           /label= Mature_human_LDCAM_polypeptide
67..69
                                                                                                   /label= Leader_peptide
39..442
                                                                                                                                                                Location/Qualifiers
                                                                                                                                         /label= Extracellular_domain
e= "N-Glycosylation
.167
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                                                             "N-Glycosylation
                                     "N-Glycosylation site"
                                                                                                                                                                                                                                                                                                                                                                         442
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Pred. No. 2.
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                                                              site"
                                                                                                                                                                                                                  modulator; immune
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?.1e-122;
                                                                                                                                                                                                                                                                                                                                                                                                                                      374
                                                                                                                                                                                                                                                                                                                                                                                                                                                              336
                                                                                                                                                                                                                                           treatment; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
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                                                                                                                                                                                                                               infectious disease;
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                                                                                                                                                                                                                                                          cell;
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ICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDALELTCEAIGKPQPVMV

ICQVEHPAVTGNLQTQRYLEVQYKPQVHTQMTYPLQGLTREGDALELTCEAIGKPQPVMV

278 240 GEEIEVNCTAMASKPATTIRWFKGNTELKGKSEVEEWSDMYTVTSQLMLKVHKEDDGVPV

GEETEVNCTAMASKPATTTRWFKGNTELKGKSEVEEWSDMYTVTSQLMLKVHKEDDGVPV

218

180

LLNFSSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVE

LLNFSSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVE

158 120 98 60

IPTGDGQNLFTKDVTVIEGEVATISCQVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQ

0;

Indels

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Gaps

0;

I PTGDGQNLFTKDVTV I EGEVAT I SCQVNKSDDSV I QLLNPNRQT I Y FRDFRPLKDSR FQ

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159 181

121 99 61 39 Qy

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219

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Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 336; Conservative 0
                                                                                    liver, spleen and heart, lung, muscle, placenta, thyroid and lung carcinoma. LDCAM polypeptides interacts with T cell surface molecules to alter signalling and inhibits T cell proliferation, bind to themselves and BTL-1, an LDCAM binding protein and increases natural killer (NK) cell populations. It may be used to measure the biological activity and as quality control reagents of LDCAM binding proteins. LDCAM may be used for treating disorders associated with malfunctioning of immune system, inflammation, autoimmune disorders, viral infected cells, infectious diseases and for killing tumour cells. They are also useful for prevention or reducing the effect of organ and bone marrow transplant rejection and for modulating T cell immune responses. LDCAM polypeptides may also be used as carriers for delivering agents attached to T cells or cells bearing B7L-1.
                                                            Sequence
                                                                                                                                                                                                                                                                                         The present amino acid sequence is the human lymphoid derived dendrtitic cell adhesion molecule, LDCAM. It is found on lymphoid derived dendrtitic cells and displays homology to adhesion molecules, B7-1 and cytoplasmic region of B7-L1. Human LDCAM is expressed in breast, retina, foetal
                                                                                                                                                                                                                                                                                                                                                                      Claim 7;
                                                                                                                                                                                                                                                                                                                                                                                                  T cell function
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Baum
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                                                                                                                                                                                                                                                                                                                                                                    Page 42-43;
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/label= Tr
396. /
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375..39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Transmembrane_domain
                                                                                                                                                                                                                                                                                                                                                                      English
Score 1761; DB 21; Pred. No. 2.1e-122; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                  are
                                                                                                                                                                                                                                                                                                                                                                                                                  capable
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RESULT 8
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                                                                                                                           Query Match
Best Local Sin
Matches 336;
                                                                                                                                                                                                                                                    component in the proliferating cell and determining modification in the level of the cell component in proliferating cell as compared with a healthy cell, where modification indicates disorder associated with TSLC1. The method is useful for detecting a cell proliferative disorder (e.g. liver, lung or pancreatic cancer) associated with tumour suppressor lung cancer 1 (TSLC1) in a subject. The invention is useful in gene therapy and for treating a cell proliferative disorder such as lung cancer (human non-small cell lung cancer), liver cancer (hepatocellular carcinoma) or pancreatic cancer associated with modification of TSLC1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detecting cell proliferative disorder associated with tumor suppressor lung cancer (TSLC) I in subject, comprises contacting proliferating cell of subject with reagent detecting TSLC1 and detecting modification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene
                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a method for detecting cell proliferative disorder associated with tumour suppressor lung cancer 1 (TSLC1) in subject. The method comprising contacting a cell component of a proliferating cell with a reagent that detects level of the cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-AUG-2000; 2000US-225264P.
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                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 49-50; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (UYJO ) UNIV JOHNS HOPKINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; hepatocellular carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human tumour suppressor lung
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Reeves
                                                                                                                                                                                                                         production, where a reagent which modulates (preferably, level in the cells, is employed. The present sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      in TSLC1 level
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .iver; lung; pancreatic cancer; cell proliferative disorder; cytostatic;
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                                                                             IPTGDGQNLFTKDVTVIEGEVATISCQVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQ 60
             LLNFSSSELKVSLTNVSISDEGRYFCOLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVE 120
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LLNFSSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVE
                                                              IPTGDGQNLFTKDVTVIEGEVATISCQVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RH, Yoshinori
                                                                                                                                           Similarity
                                                                                                                                                                                            442 AA;
                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first
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                                                                                                                                          100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SCHOOL MEDICINE
                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer 1 (TSLC1) polypeptide
                                                                                                                                          Score 1761; DB 23; Pred. No. 2.1e-122;
                                                                                                                            Mismatches
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                                                                                                                            0
                                                                                                                            Indels
                                                                                                                                                       Length
                                                                                                                                                                                                                           human TSLC1
                                                                                                                                                                                                                                          increases)
                                                                                                                                                            442;
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                                                              98
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RESULT 9
AAY45093
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                                                                                                                                                                                                                                                                                                                                                                                                            Lymphoid derived dendritic cell adhesion molecule; LDCAM; mouse; B7-1; B7-L1; T cell proliferation; natural killer cell; NK; tumour cell; biological activity; quality control reagent; treatment; inflammation; immune system disorder; autoimmune; viral infection; infectious disease; organ transplant rejection; bone marrow; modulator; immune response.
                                      WPI; 2000-205712/18
N-PSDB; AAZ50883.
                                                                                                                              05-AUG-1999;
                                                                                                                                                                                                  Domain
                                                                                                                                                                                                                                                                                                                                                                       Key
                                                                                                                                                                                                                                                                                                                                                                                          Mus sp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse lymphoid derived dendritic cell adhesion molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-MAY-2000
                  Novel molecules designated LDCAM are capable of altering or modulating
                                                                   Baum PR,
                                                                                      (IMMV ) IMMUNEX CORP
                                                                                                           07-AUG-1998;
                                                                                                                                                                    WO200008158-A2
                                                                                                                                                                                                                     Domain
                                                                                                                                                                                                                                                                                                   Modified-site
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                                                                                                                                                                                                                                                                                                                                                              Domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAY45093 standard;
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         cell function
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49..51
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357..377
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290..292
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286..288
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147..149
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95..97
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83..85
                                                                                                                                                                                                  /label- Transmembrane_domain 378..423
                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Protein;
                                                                                                                                                                                                                                                                                        "N-Glycosylation site"
                                                                                                                                                                                                                                                                                                           "N-Glycosylation site"
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RESULT 10
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Best 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         region of B7-LI. Mouse LDCAM is found on whole embryo, testes, triple cells murine splenic and lymph node CD8+, S49.1 and dendritic cells. LDCAM polypeptides interacts with T cell surface molecules to alter signalling and inhibits T cell proliferation, bind to themselves and B7L-1, an LDCAM binding protein and increases natural killer (NK) cell populations. It may be used to measure the biological activity and as quality control reagents of LDCAM binding proteins. LDCAM may be used for treating disorders associated with malfunctioning of immune system, inflammation, autoimmune disorders, viral infected cells, infectious diseases and for killing tumour cells. They are also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cell adhes
                                                                         Crohn's disease; nephriti cardiovascular disorder;
                                                                                                                      anticancer; vulnerary; antiviral; antibacterial; antifungal; immune disorder; Addison's disease; rheumatoid arthritis; de
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  useful for prevention or reducing the effect of organ and bone marrow transplant rejection and for modulating T cell immune responses. LDCAM polypeptides may also be used as carriers for delivering agents attached to T cells or cells bearing B7L-1.
                                                          melanoma;
                                                                                                       multiple sclerosis; inflammatory disorder; inflammatory bowel disease;
                                                                                                                                                       antirheumatic, dermatological; antiproliferative; antiarteriosclerotic;
                                                                                                                                                                       Secreted protein; immunosuppressant; anti-inflammatory;
                                                                                                                                                                                                                                          21-NOV-2000
                                                                                                                                                                                                                                                                                                       AAB25586
                                                                                                                                                                                                     Protein encoded by human secreted protein gene #11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                      321
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IPTGDGQNLFTKDVTVIEGEVATISCQVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDAFELTCEAIGKPQPVMV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDALELTCEAIGKPQPVMV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LLNFSSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cells or cells bearing B7L-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       332;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page 46-47; 44pp; English
                                                          lymphoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nt amino acid sequence is the mouse lymphoid derived dendrtitic sion molecule, LDCAM. It is found on lymphoid derived dendrtitic displays homology to adhesion molecules, B7-1 and cytoplasmic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                       (first entry)
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                                                                                        nephritis; hyperproliferative disorder;
                                                                                                                                                                                                                                                                                                    Protein;
                                                           wound
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                                                                          coronary arteriosclerosis;
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Pred. No. 5
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                                                           numan
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5.9e-121;
hes 3;
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                                                                          myocarditis;
                                                                                                                                                                         antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     423;
                                                                                                                         dermatitis;
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                                                                          cancer
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339 301 279 241 219

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TWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIP

TWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIP

ICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDALELTCEAIGKPQPVMV

ICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDALELTCEAIGKPQPVMV

278

GEEIEVNCTAMASKPATTIRWFKGNTELKGKSEVEEWSDMYTVTSQLMLKVHKEDDGVPV

GEETEVNCTAMASKPATTIRWFKGNTELKGKSEVEEWSDMYTVTSQLMLKVHKEDDGVPV

218 180 158 120 98 60

LLNESSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTTTVLVPPRNLMIDIQKDTAVE

LLNFSSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVE

Conservative

Indels

0;

Gaps

0

121

99 61 39

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Matches
                   Query Match
Best Local
                                                                                                                                                                                                                                    in AAA80606-A80623 encode the 12 secreted protein sequences given in AAB2576-B2593. The human secreted proteins have various activities dependent on the tissues in which they are expressed. Examples of the activitles of the proteins include: immunosuppressant; anti-inflammatory; antiarthritic; antirheumatic, dermatological; antiproliferative; antiarteriosclerotic; anticancer; vulnerary; antivital; antibacterial; and antifungal activity. The proteins, polypeptides, agonists and antagonists may be used to treat prevent and/or diagnose various disease, disorders and conditions examples of which include: immune disorders e.g. Addison's disease, rheumatoid arthritis, dermatitis, and multiple sclerosis; inflammatory disorders e.g. Inflammatory bowel disease, Crohn's disease and nephritis; e.g. Malaneary bowel disease, Crohn's disease and nephritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 200
N-PSDB;
                                                                                                                      hyperproliferative disorders such as paraproteinaemias and purpura; cardiovascular disorders e.g. coronary arteriosclerosis and myocarditis; cancer e.g. melanoma and lymphoma. The proteins and polynucleotide sequences may also be used in wound healing and the treatment of infectious diseases. The human secreted protein gene #11 and protein sequences are represented in sequences AAABOSOB6. Sequences sequences are represented in sequences AAABOSOB6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolated nucleic acid molecules encoding human secreted proteins are used for the prevention, amelioration and treatment of autoimmune, inflammatory, hyperproliferative and cardiovascular disorders, cancewounds, and infectious diseases.
                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleotide sequences encoding them. The point AAA80606-A80623 encode the 12 secreted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Figure 28A-B; 803pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Greene
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                                                                                                              AAA80677-A80682 represent
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                   Local Similarity
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DB; AAA80616.
325;
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                                                                         364
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention relates to 12 secreted human proteins and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SM,
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                 96.8%;
100.0%;
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                                                                                                            genes related to
0,
                 Score 1704; DB 2
Pred. No. 2.7e-1
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Mismatches
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                                                                                                            the secreted protein
                                  21;
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                                    Length
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                                    364;
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                                                                                                            gene#11
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RESULT 11
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Matches 312
                                                                                                                                                                                                             can be used in gene therapy. The polynucleotide sequences and the proteins they encode may be used in the prevention, treatment and diagnosis of diseases associated with inappropriate secretory protein/membrane protein expression. The nucleic acids and complementary sequences may also be used as DNA probes in diagnostic assays (e.g. polymerase chain reactions (PCR)) to detect and quantitate the presence of similar nucleic acid sequences in samples. They may also be used to study the expression and function of secretory proteins/membrane polypeptides and their role in metabolism. The polypeptides may be used as antigens in the production of antibodies against them and in assays to identify modulators (agonists and antagonists) of expression and activity. The antibodies and antagonists may also be used as therapeutic agents to down regulate expression and activity. The antibodies may also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-JUL-1999; 99JP-0194179
11-JAN-2000; 2000JP-0118775
02-MAY-2000; 2000JP-0183766
                                                                                                                                                  polypeptides in samples (e.g. (ELISA). Examples of diseases arthritis and diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                  cDNA sequences of the invention. The invention also includes methods for the production of antibodies directed against the proteins, and cDNA sequences, which can be used in vaccines. The polynucleotide sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acids encoding gene therapy or as cand
                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAB88317 - AAB88419. Included in the invention are primers AAF93917 - AAF94295 and AAF62232 - AAF62235 which are used to isolate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention relates to nucleic acid sequences AAF93744 - AAF93916 which encode human secretory or membrane proteins represented by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-JUL-2000; 2000EP-0114090
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                                                                                                                      Sequence
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 39
                                                                                                                                                                     used as diagnostic agents for detecting the presence of the lypeptides in samples (e.g. by enzyme linked immunosorbant assay LISA). Examples of diseases which may be treated include rheumat
                              μ
2001-093989/11.
                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard;
                                                                                                                          443
                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ncoding secretory proteins/membrane pro
as candidate target molecules in drug
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                                                                           91.6%;
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                                                           Score 1612.5;
Pred. No. 2.1e
4; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English.
                                                                             . 1e-1
                                                                                                                                                                      treated include rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sugiyama
                                                      DB 2
1111;
20;
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                                                           Indels
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                                                                                        Length 443;
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                                                         Gaps
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RESULT 12
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02-JUN-1998;
22-JUL-1998;
31-JUL-1998;
10-AUG-1998;
11-AUG-1998;
                                                                                                                                                                                                                                                                                                                                              numan; secreted protein; nutritional; cytokine; cell proliferation; differentiation; immune stimulating; vaccine; suppression; haematopoiesis regulation; tissue growth; activin; inhibin; chemotactic; chemokinetic; haemostatic; thrombolytic; receptor; ligand; anti-inflammatory; cadherin; tumour invasion suppressor; tumour inhibition; gene therapy.
                                                                                               WPI; 200
N-PSDB;
                                                                                                                                       Jacobs K
Merberg
                                                                                                                                                                    (GEMY
                                               Claim
                                                                                                                              DiBlasio-Smith
                                                                                                                                                                                        06-MAY-1999
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                                                                           polynucleotides
                                                                                               2000-052937/04.
DB; AAZ33346.
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                                                                                                                                   placenta,
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                                              Page 416-417;
                                                                                                                           McCoy JM,
Treacy M,
mith E, W
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98US-0087645.
98US-0093712.
98US-0094935.
98US-0095880.
98US-0096068.
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                                                                 ides encoding secreted huma adult retina, fetal brain,
                                                                                                                            Widom
                                                                                                                                                Lavallie
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                                              English
                                                                                                                                       Collins-Racie L
Steininger RJ,
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The present invention describes new human secreted proteins which were isolated from adult placenta, adult retina, foetal brain, foetal kidney, adult blood, adult brain, adult thyroid, adult blood, adult neural tissue, adult testes, and adult lymph node cDNA libraries. The human

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RESULT 1:
AABZ5593
ID AABZ5593
AC AABS
XX AABC2593
AC AABC
XX Prot
DT 21-b
CXX Secr
KW Secr
KW antit
KW antimul
KW inmu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              secreted proteins, and the polynucleotides encoding them, are predicte to have biological activities which would make them suitable for treating, preventing or ameliorating medical conditions in humans and animals. Suggested activities include nutritional activity, cytokine and cell proliferation/differentiation activity, immune stimulating (e.g. as vaccines) or suppressing activity, haematopoiesis regulating activity, tissue growth activity, activin/inhibin activity, chemotactic/chemokinetic activity, haemostatic and thrombolytic activity, receptor/ligand activity, anti-inflammatory activity, cadherin/tumour invasion suppressor activity, and tumour inhibition activity. The polynucleotides are also stated to be useful for gene therapy. AAZ33316 to AAZ33373 encode human secreted proteins, and AXY52998 to AAX53060 represent human secreted proteins, given in the
                 27-OCT-1999;
                                                                                                                                                                 Crohn's disease; nephritis; hyperproliferative disorder; cardiovascular disorder; coronary arteriosclerosis; myoc
                                                  25-MAY-2000
                                                                                                                                                   melanoma;
                                                                                                                                                                                                                                                                Secreted protein; immunosuppressant; anti-inflammatory;
                                                                                                                                                                                                                                                                                                    Protein encoded
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                                                                                 WO200029435-A1
                                                                                                                    Homo
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                 99WO-US25031
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Pred. No. 7.5e-110;
0; Mismatches 0;
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                                                                                                                                               human.
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AAM78418

standard;

Protein;

387

Human; Human 06-NOV-2001

protein cytokine;
e; peptide

SEQ ID NO 1080.

cell proliferation;
therapy; stem cell ;

growth

cell differentiation;

gene

therapy

(first entry)

Qy

В

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181 290 121

LYVYDPPTTIPPPTTTTTTTTTTTTTLTIITDSPSQVKKARSGQ

LYVYDPPTTIPPPTTTTTTTTTTTTILTIITDS-----

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EATGKPQPVMVTWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYM

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                                                                                                                                                                                                                                                                                                                                                                                     CC AAB35576-B2593. The human secreted proteins have various activities of the tissues in which they are expressed. Examples of the CC activities of the proteins include: immunosuppressant; antiplation; antiarthritic; antirheumatic, dermatological; CC antiproliferative; antiarteriosclerotic; anticancer; vulnerary; CC antiproliferative; and antisponists may be used to treat prevent and/or diagnose various disease, disorders and conditions examples of cond/or diagnose various disease, disorders and conditions examples of cond/or diagnose various disease, disorders and conditions examples of conditions examples of conditions examples of conditions of the conditions of th
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                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Isolated nucleic acid molecules encoding human secreted proteins are used for the prevention, amelioration and treatment of autoimmune, inflammatory, hyperproliferative and cardiovascular disorders, cance:
                                                                                                                                                                                                                                                                                                                                                                                                                                AAAB0677-A80682 represent genes related to the secreted protein gene#11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      inflammatory, hyperproliferative wounds, and infectious diseases
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EAIGKPQPYMYTWVRVDDEMPQHAYLSGPNLFINNLNKTDNGTYRCEASNIYGKAHSDYM
                                                                                                     KVHKEDDGVPVICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDALELTC
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                                                                                                                                                                                                                                                                                                                                                                                     229 AA;
                                                                                                                                                                                                                                                                                Conservative
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Pred. No. 2.4e
2; Mismatches
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27-APR-2000;
20-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to polynucleotides (AAK51456-AAK53435) and the encoded polypeptides (AAM78323-AAM80302) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polynucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, hammatopolesis regulating activity, tissue growth factor activity, hammatopolesis regulating activity, tissue growth factor activity, the consideration activity and may be useful in the diagnosis and/or treatment of cancer, leukaemia, nervous system disorders, arthritis and
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Zhao
                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 20; Page 3307-3308; 6221pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acids encoding polypeptides with cytokine-like useful in diagnosis and gene therapy -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N-PSDB;
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01-SEP-2000;
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20-OCT-2000;
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          WVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIPP
                                       AQNQALPSGHSKQTQYVLDVQYSPTARIHAS---QAVVREGDTLVLTCAVTGNPRPNQIR
                                                                 VEHPAVTG - - NLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDALELTCEAIGKPQPVMVT
                                                                                            VELSCLVPRSRPAATLRWYRDRKELKGVSSSQENGKVWSVASTVRFRVDRKDDGGIIICE
                                                                                                                     IEVNCTAMASKPATTIRWFKGNTELKGKSEVEEWSDMYTVTSQLMLKVHKEDDGVPVICQ 183
                                                                                                                                                                            FSSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVEGEE 123
                                                                                                                                                  FSPRRVRIRLSDARLEDEGGYFCQLYTEDTHHQIATLTVLVAPENPVVEV-REQAVEGGE
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, Wang D,
Yang Y,
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Wejhrman T,
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Wang J,
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, Zhang J, Ren
n T, Goodrich R
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Pred. No. 1.3e-38;
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R, Wang
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            Query Match
Best Local :
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                                                                This sequence is the human beta-secretase enzyme polynucleotide. Beta-secretase is capable of cleaving the beta-amyloid precursor protein (APP) (AAV33742;swedish mutant APP). This enzyme is used in a method for detecting human beta-secretase cleavage of polypeptides and for identifying beta-secretase inhibitors. The protein has a deduced molecular weight of 49 kD, although this is without glycosylation and the expected molecular weight is therefore higher. Inhibition of beta-secretase activity would be useful for chemical modelling of a critical event in the pathology of Alzheimer's disease. Inhibitors of beta-secretase would be useful for the prevention and treatment of Alzheimer's disease and Down's Syndrome.
                                                                                                                                                                                                                                                                                                                                                        07-JUN-1996;
07-JUN-1995;
07-JUN-1995;
                                              Sequence
                                                                                                                                                                                                                                      A method for detecting useful for identifying
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                                                                                                                                                                                                                                                                                                           JP,
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                                                444 AA;
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95US-0485152.
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24..444
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            35.48;
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GONLFTKDVTVIEGEVATISCQVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQLLNFS

GQFPLTQNVTVVEGGTATLTCRVDQNDNTSLQWSNPAQQTLYFDDKKALRDNRTELVRAS

Search completed: November 20, 2002, 07:37:56 Job time: 65.5 secs

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Title:
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1: sp_archea:*
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7: sp_mhc:*
9: sp_phage:*
10: sp_plant:*
11: sp_vortebrate:*
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13: sp_vertebrate:
14: sp_unclassifie
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Gapop 10.0 , Gapext 0.5
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1761
1 IPTGDGQNLFTKDYTVIEGE.....LTIITDSRAGEEGSIRAVDH 336
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Copyright (c) 1993 - 2002 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

16	15	14	13	12	11	10	9	8	7	6	G	4	ω	N	-	Result
298	301	469.5	482.5	609	622	1024.5	1048	1137	1157.5	1180	1529	1725.5	1725.5	1741	1758	Score
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09ЛГВ9	Q9D006	Q9UJP1	Q99N28	Q9Y4A4	Q8R464	Q9QYL3	Q9QYL5	09QYL6	Q9QYL4	Q9Z2H8	Q9D6E7	Q9CRY3	Q8R5M8	Q8R4L1	Q9BY67	ID
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216	216.5	216.5	217	217.5	217.5	218	219	219	220	220	221	222	222	222	223	223.5	223.5	224.5	227	235	243	243	248	250	263.5	277.5	277.5	292
12.3	12.3	12.3	12.3	12.4	12.4	12.4	12.4	12.4	12.5	12.5	12.5	12.6	12.6	12.6	12.7	12.7	12.7	12.7	12.9	13.3	13.8	13.8	14.1	14.2	15.0	15.8	15.8	16.6
392	1270	975	417	968	959	1032	1060	344	449	344	344	1675	1380	467	725	34350	7962	605	4162	393	1482	725	407	439	5198	510	438	549
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Matches
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FUKAMI T., Maruyama T., Murakami Y.;
"Identification of murine orthologue of the TS
Submitted (OCT-2001) to the EMBL/GenBank/DDBJ
EMBL; AF434663; AAL86736.1; -
SEOUENCE 445 AA; 48664 MW; C5D5A070DAF70E!
                                                                                                                                                                                                                                                                                                                                                                             Q8R4L1;
Q8R4L1;
Q1-JUN-2002
Q1-JUN-2002
Q1-JUN-2002
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Mammalia; Eutheria;
NCBI_TaxID=10090;
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                                                                                               GEEIEVNCTAMASKPATTIRWFKGNTELKGKSEVEEWSDMYTVTSQLMLKVHKEDDGVPV
                                                                TWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIP
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                                                                                                                                                                                                                                            Similarity
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(TremBLrel. 21, Last sequence up
(TremBLrel. 21, Last annotation
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Rodentia;
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; Murinae; Mus
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Q9CRY3;
Q9CRY3;
Ol-JUN-2001 (TrEMBLrel. 17, C
Ol-JUN-2001 (TrEMBLrel. 17, L
Ol-DEC-2001 (TrEMBLrel. 19, L
3100001108Rik protein (Fragme
IGSF4 OR 3100001108RIX.
Mus musculus (Mouse)
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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Eukaryota; Metazoa; (Mouse)
Mammalia; Eutheria; (Mouse)
NCBI_TaxID=10090;
SEQUENCE FROM N.A.
STRAIN-C57BL/6J; TISSUB-EMBRYONIC HEAD;
MEDLINE-2108560; PubMed-11217851;
MEDLINE-2108560; PubMed-11217851;
Kawai J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ish
Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fuku
Arakawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamana
Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Sai
Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant
Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H
Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quacke
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01-JUN-2002
01-JUN-2002
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Momoi T.;
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Rodentia;
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1. 19, Last annotation update)
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Pred. No. 3.7e
1; Mismatches
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Sciurognathi;
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thi; Muridae;
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                            oh M., Ishii
i J., Fukuda
S., Yamanaka
va T., Saito
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; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
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; Murinae; Mus
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  chiwa H.,
Quackenbush
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RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,

RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,

RA Brownstein M.J., Bult C., Fletcher C., Fijita M., Gariboldi M.,

RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,

RA Lyons P., Marchionni L., Mashima J., Mazzarelli J., Mombaerts P.,

RA Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,

RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,

RA Sasaki H., Toyo-oka K., Wang K.H., Weltz C., Whittaker C., Wilming L.,

RA Wynshaw-Boris A., Yoshida K.H., Weltz C., Whittaker C., Wilming L.,

RA Hayashizaki Y.,

RT "Functional annotation of a full-length mouse cDNA collection.";

RL Nature 409:685-690(2001).

RE EMBL: AKO13911: BAB29050.1; -.
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Best Local Sim
Matches 332;
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Q9D6E7;
Q9D6E7;
Q1-JUN-2001 (TrEMBLrel. 17, Cre
Q1-JUN-2001 (TrEMBLrel. 17, Las
Q1-DEC-2001 (TrEMBLrel. 19, Las
Q1-DEC-2001 (TrEMBLrel. 19, Las
Q100073G66R1k protein.
IGSF4 OR 2900073G6R1K.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; C
Mammalia; Eutheria; Rodentia; S
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InterPro: IPR003508; Ig.2.
InterPro: IPR003500; Ig_NHC.
InterPro: IPR003006; Ig_MHC.
Pfam: PF00047; Ig: 3.
SMART: SM00409; IG: 3.
SMART: SM00408; IG: 2; 3.
SMART: SM00410; IG.11ke; 1.
Immunoglobulin domain.
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SEQUENCE FROM N.A.
STRAIN=C57BL/6J; TISSUE=HIPPOCAMPUS;
MEDLINE=21085660; PubMed=11217851;
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                                                                                            NCBI_TaxID-10090;
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hes 332; Conservative
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Pred. No. 4.1e-136;
1; Mismatches 3;
                                                                                                                Craniata; Ver
Sciurognathi;
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thi; Muridae; Murinae; Mus
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RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Salto T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Salto R.,
RA Salto T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Salto R.,
RA Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Sakai K., Okido T., Fletcher C., Fujita M., Gariboldi M.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
RA Lyons P., Marchionni L., Mashima J., Mazazarelli J., Mombaerts P.,
RA Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,
RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,
Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Kohtsuki S.,
RA Hayashtaki Y.;
RA Hayashtaki Y.;
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Best Local :
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0922H8;
01-MAY-1999 (TrEMBLrel. 1
01-MAY-1999 (TREMBLrel. 1
01-JUN-2002 (TREMBLrel. 2
Nectin 11ke protein 2.
IGSF4 OR NECL2.
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SMART; SM00409; IG; 3.
SMART; SM00408; IG; 3.
SMART; SM00410; IG_like; 1.
Immunoglobulin domain.
SEQUENCE 336 AA; 37157 MW;
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InterPro;
InterPro;
                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=10090;
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EMBL; AK013775; BAB28988.1; -.
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                                                                              Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     37157 MW;
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99.3%;
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Last annotation updat
                                                                                                                                                                                                                                                                     Created)
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Pred. No. 6
                                                                           Craniata; Ver
Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FF887FAF4EFDF120 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                          Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 11;
5.7e-120;
                                                                                                                                                                                                                                                                                                                          B
                                                                                 Muridae;
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RESULT OPOCILIA OPOCI
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Query Match 65.7
Best Local Similarity 93.7
Matches 222; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                           "RA175, a novel neuron specific submitted (DEC-1998) to the EMBI EMBL, AB021966; BAA87916.1; -MGD; MGI:1889272; Igsf4. InterPro; IPR003598; Ig_C2. InterPro; IPR003600; Ig_Nike. InterPro; IPR003006; Ig_MHC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro: IPR003585; Neurexin-like.
Pfam; PF00047; ig; 2.
SMART; SM00294; 4.lm; 1.
SMART; SM00408; IGC2; 1.
SMART; SM00410; IGC2; 1.
Immunoglobulin domain.
Immunoglobulin domain.
SEQUENCE 295 AA; 32509 MW; 9DE9DE
                                                                                                                                                                                                                        SMART;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAY-2000 (TrEMBLrel. 13,
01-MAY-2000 (TrEMBLrel. 13,
01-JUN-2002 (TrEMBLrel. 21,
Adhesion protein RAI/5C.
IGSF4 OR RAI/5C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zhou Y., Du G., Chen J., Yuan J., Qiang B.;
"Cloning of a novel cDNA encoding a member of
Submitted (APR-1988) to the EMBL/GenBank/DDBJ
EMBL; AF061260; AAC67243.1; -.
MCD; MCI:1889272; Igsf4.
InterPro; IPR003598; Ig_C2.
InterPro; IPR003506; Ig_Like.
InterPro; IPR003006; Ig_MHC.
                                                                                                                                                                                                                                                                                          Pfam;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Momoi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A. Soyama A., Fujita
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse) Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                           InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         110 MIDIQKDTAVEGEEIEVNCTAMASKPATTIRWFKGNTELKGKSEVEEWSDMYTVTSQLML
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                                                                                                                                                                                 N; PF00047; 19; 2.
AT; SM00294; 4.1m; 1.
AT; SM00408; IGC2; 1.
AT; SM00410; IG_like; 1.
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                                                                                                                                                                                                                                                                                                                       IPR003585; Neurexin-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    s (Mouse).

Metazoa; Chordata; C
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                                                                                                                                                             domain
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                                65.7%;
93.3%;
                                                                                                                            33522 MW;
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Last sequence update)
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Score 1157.5; DB 11; Length Pred. No. 7.3e-89; 2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRT;
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3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kouroku
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               databases
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                                                                                                                            CRC64;
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ID Q9QYL5
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Matches
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                                                                                                                                                                                                                                                                                         "RA175, a novel neuron specific adhesion pr
Submitted (DEC-1998) to the EMBL/GenBank/DD
EMBL; AB021964; BAA87914.1; -.
MGD; MGI:1889272; Igsf4.
InterPro; IPR003598; Ig_c2.
InterPro; IPR003506; Ig_like.
InterPro; IPR003006; Ig_Hike.
InterPro; IPR003006; Ig_Hike.
InterPro; IPR003585; Neurexin-like.
Pfam; PF00047; ig; 2.
SMART; SM00249; 4.1m; 1.
SMART; SM00410; IG_like; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IGSF4 OR RAI....
Mus musculus (Mouse).
Mus musculus (Mouse).
Portaryota; Metazoa; Chordata;
Portaryota; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-MAY-2000 (TrEMBLrel.
01-MAY-2000 (TrEMBLrel.
01-JUN-2002 (TrEMBLrel.
Adhesion protein RA175A.
                                                                                                                                                                                                                                                                        SEQUENCE
                                                                                                                                                                                                                                                                                    Immunoglobulin
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Soyama A., Fujita E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Adhesion protein IGSF4 OR RA175A.
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mes 215; Conserv
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                                                                 LYVYDPPTTIPPPTTTTTTTTTTTTTTILTIITDSRAGEEGSIRAVDH
                                                       LYVYDPPTTIPPPTTTTTTTTTTTTTEPAVHDSRAGEEGTIGAVDH
                                                                                                EAIGKPQPVMVTWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYPCEASNIVGKAHSDYI
                                                                                                           EAIGKPQPVMVTWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYM
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                                                                                                                                                                                                                               Conservative
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94.7%;
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EMBL/GenBank/DDBJ databases.
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Pred. No. 3.6e
3; Mismatches
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Sciurognathi; Muridae;
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annotation
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; Murinae; Mus
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Best Local Sin
Matches 203;
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InterPro; IPR003600; Ig_11ke.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003085; Neurexin-like.
InterPro; IPR003585; Neurexin-like.
Pfam; PF00047; 19; 2
SMART; SM00294; 4.1m; 1.
SMART; SM00400; IGC2; 1.
SMART; SM00410; IG_11ke; 1.
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O90YL3
O1-MAY-2000 (TrEMBLrel. 1
O1-MAY-2000 (TrEMBLrel. 1
O1-JUN-2002 (TrEMBLrel. 2
O1-JUN-2002 (TrEMBLrel. 2
Adhesion protein RA175N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted
Soyama A., Fujita E., Urase K., Mukasa T., Kouroku Momol T.;
"RA175, a novel neuron specific adhesion protein.";
submitted (DEC-1998) to the EMBL/GenBank/DDBJ datab EMBL; AB021967; BAA87917.1; -...
MGD; MGT:1889272; Igsf4.
InterPro; IPR003598; Ig_C2.
InterPro; IPR003600; Ig_like.
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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01-MAY-2000 (TrEMBLrel.
01-MAY-2000 (TrEMBLrel.
01-JUN-2002 (TrEMBLrel.
                                                                                               SEQUENCE FROM
                                                                                                                              Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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Sciurognathi; Muridae;
                                                                                                                              Craniata; Vertebrata; | Sciurognathi; Muridae;
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01-JUN-2002 (TrEMBLrel. 21
01-JUN-2002 (TrEMBLrel. 2:
01-JUN-2002 (TrEMBLrel. 2:
Membrane glycoprotein.
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Best Local
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InterPro; IPR003585; Neurexin
Pfam; PF00047; 1g; 2.
SMART; SM00294; 4.ln; 1.
SMART; SM00294; 4.ln; 1.
SMART; SM00408; IGC2; 1.
SMART; SM00410; IG_Like; 1.
Immunoglobulin domain.
SEQUENCE 278 AA; 30636 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Identification of a murine ortholog of the TS Submitted (OCT-2001) to the EMBL/GenBank/DDBJ EMBL; AY059394; AAL29692.1; ... SEQUENCE 388 AA; 42723 MW; 8E3A9DF1C3B9D2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalla; Eutherla; Rodentia;
NCBI_TaxID=10090;
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STRAIN-129/SVJ;
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242
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                                                                                 VEHPAVTG--NLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDALELTCEAIGKPQPVMVT
                                                                                                                                    FSSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVEGEE 123
                                                                                                                                                                                                                                                                                                                                                GDGQNLFTKDVTVIEGEVATISCQVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQLLN
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WVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIPP
                                      AQNQALPSGHSKQTQYVLDVQYSPTARIHAS---QAVVREGDTLVLTCAVTGNPRPNQIR
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Pred. No. 8.7e
3; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. 6.6e-
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8E3A9DF1C3B9D23E CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       622;
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                                                                                                                                                                                                                                                                                                                                                                                        DB 1
6.6e-44;
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A Burkhart-Schultz K.J., Gordon L., Kyle A., Ramirez M., Stilwagen
A Phan H., Velasco N., Do L., Regala W., Terry A., Garnes J.,
A Danganan L., Poundstone P., Christensen M., Georgescu A., Avila
A Liu S., Attix C., Andreise T., Trankheim M., Amico-Keller G.,
Coefield J., Duarte S., Lucas S., Bruce R., Thomas P., Quan G.,
Coefield J., Duarte S., Lucas S., Bruce R., Thomas P., Quan G.,
A Kronmiller B., Arellano A., Montgomery M., OW D., Nolan M., Tron
A Kobayashi A., Olsen A.S., Carrano A.V.;
Tsequence analysis of a 2.5 Mb region in 19q13.2 containing a
T clustered CEA/PSG gene family.";
Submitted (AUG-1998) to the EMBL/GenBank/DDBJ databases.
R EMBL, ACOS525; AAC32740.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pfam; PF00047; ig; 3.
SMART; SM00294; 4.1m; 1.
SMART; SM00408; IGc2; 1.
SMART; SM00410; IG_like;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Q9Y4A4
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                                                                                                                                                                                                                                                                                                           VNCTAMASKPATTIRWFKGNTELKGKSEVEEWSDMYTVTSQLMLKVHKEDDGVPVICQVE
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                    ---IPPPTTTTTTTTTTTL
                                                                                                                                                                                                                         HPAVTG---NLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDALELTCEAIGKPQPVMVTWV
                                                                                                                                                                                                                                                                                                                                                                               PRRVRIRLSDARLEDEGGYFCQLYTEDTHHQIATLTVLVAPENPVVEV-REQAVEGGEVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GQEVQTENVTVAEGGVAEITCRLHQYDGSIVVIQNPARQTLFFNGTRALKDERFQLEEFS
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                                                                        RGNESLPERAEAVGETLTLPGLVSADNGTYTCEASNKHGHARALYVLVVYGESRLRPTEG
                                                                                                                      RVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDP---
                                                                                                                                                                          NQALPSGHSKQTQYVLDVQYSPTARIHAS---QAVVREGDTLVLTCAVTGNPRPNQIRWN
                                                                                                                                                                                                                                                                          LSCLVPRSRPAATLRWYRDRKELKGVSSSQENGKVWSVASTVRFRVDRKDDGGIIICEAQ
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; IPR003600; Ig_like.
; IPR003006; Ig_MHC.
; IPR003585; Neurexin-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         304
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Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           34.6%; Score 609; DE 39.1%; Pred. No. 7.96 tive 62; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Neurexin-like.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Last sequence up
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 4; Length .9e-43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AA
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Best Local
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01-JUN-2001 (
01-JUN-2002 (
Nectin-like F
                                                                                                                                                                                                                                                                                                                                                                                       InterPro;
InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                       "Identification of a murine ortholog of the TSLC1-like gene Submitted (OCT-2001) to the EMBL/GenBank/DDBJ databases. EMBL; AF195662; AAG35584.1; -. EMBL; AY059393; AAL29691.1; -. MGD; MGI:2137858; Necll.
                                                                                                                                                                                                                                                                                                                                SMART; SM00409; IG; 3.
SMART; SM00408; IGc2; 3.
SMART; SM00410; IG_like;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Zhou Y., Huang X., Yuan "Cloning and expression membrane protein.";
Submitted (OCT-1999) to
                                                                                                                                                                                                                                                                                                               SEQUENCE
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                                                                                                                                                                                                                                                                                                                         Immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                  Pfam; PF00047; ig; 3
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InterPro; IPR003006;
                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR003599;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fukami T., Maruyama T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN=129/SVJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=10090;
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 319
                     297
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                                                                                                  CQVEHPAVTG-NLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDALELTCEAIGKPQPVMV
                                                                                                                                               EVNCTAMASKPATTIRWFKGNTELKG-KSEVEEWSD--MYTVTSQLMLKVHKEDDGVPVI
                                                                                                                                                                             TPHELSISISNVALADEGEYTCSIFTMPVRTAKSLVTVLGIPQKPIITGYKSSLREKETA
 --- PVPSSSST
                    TTIPPPTTTT
                                          VWVKEGSEPPLKMTQESALIFP---
                                                                TWVRVDDEMP----QHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPP
                                                                                                                                 TLNCQSSGSKPAAQLTWRKGDQELHGDQTRIQEDPNGKTFTVSSSVSFQVTREDDGANIV
                                                                                                                                                                                                  SSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVEGEEI 124
                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                          IPR003598; Ig_c2.
IPR003600; Ig_like.
IPR003006; Ig_MHC.
                                                                                                                                                                                                                                                                                                               396 AA;
                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (TrEMBLrel. 17, Last sequence upd:
(TrEMBLrel. 21, Last annotation uprotein 1 (Membrane glycoprotein)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (Mouse)
                                                                                                                                                                                                                                                                                                                         domain
 326
                     307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                            42964 MW;
                                                                                                                                                                                                                                                                            27.48;
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17,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              J., Qiang B.
analysis of
                                                                                                                                                                                                                                                                   63;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Created)
                                                                                                                                                                                                                                                                            Score 482.5; DB 1
Pred. No. 3.2e-32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Craniata; Veri
Sciurognathi;
                                          ---FLNKSDSGTYGCTATSNMGSYTAYFTLNVNDPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRT;
                                                                                                                                                                                                                                                                                                              C1ADF8B57D141F3A CRC64;
                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                novel
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                                                                                      PEPAHPREGQKLLLHCEGRGNPVPQQY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vertebrata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ΑA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mouse cDNA encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     update)
                                                                                                                                                                                                                                                                                       DB 11; Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Muridae;
                                                                                                                                                                                                                                                                    119;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Murinae;
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RESULT Q9UJP1

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RESULT 15
Q9D006
ID Q9D00
AC Q9D00
DT 01-JU
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Best Local
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InterPro: IPR003600; Ig_like.
InterPro: IPR003006: Ig_like.
InterPro: IPR003006: Ig_HMC.
InterPro: IPR003585; Neurexin-like.
Pfam; PF00047; Ig; 3.
SMART: SM00294; 4.1m; 1.
SMART: SM002408; IGc2; 1.
SMART: SM00410; IG_like; 2.
                                      Q9D006 PRELIMINARY;
Q9D006;
01-JUN-2001 (TrEMBLrel. 1
01-JUN-2001 (TrEMBLrel. 1
01-DEC-2001 (TrEMBLrel. 1
2610301B19R1k protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zhou
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
BK134P22.1 (Novel protein similar to mouse IMMUNOSUPERFAMILY protein
BL2) (NECTIN-11ke protein 1).
BK134P22.1 OR NECL1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q9UJP1;
             Mus musculus (Mouse).
                             2610301B19RIK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Zhou Y., Du G., Yuan J., Olang
Submitted (APR-2001) to the EM
EMBL; AL035403; CAB56227 1; -.
EMBL; AF062733; AAD17540.2; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID-9606;
                                                                                                                                                                           361
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                                                                                                                                                                                                                                                                                                                                                                                                           KVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVEGEEIEVNCT 129
                                                                                                                                                                            :- ES
                                                                                                                                                                                                                                                         DDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIPPPTTT
                                                                                                                                                                                                                                                                                                                                                     AMASKPATTIRWFKGNTELKGK-SEVEEWSD--MYTVTSQLMLKVHKEDDGVPVICQVEH 186
                                                                                                                                                                                                                                                                                                                                                                                          SISISNVALADEGEYTCSIFTMPVRTAKSLVTVLGIPQKPIITGYKSSLREKDTATLNCQ
                                                                                                                                                                                                                                                                                                                                                                                                                                              WTSDETVVAGGTVVLKCQVKDHEDSSLQWSNPAQQTLYFGEKRALRDNRIQLVTSTPHEL 127
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                                                                                                                                                                                                                                                                                  ESLKGADRSTSQRIEVLYTPTAMIRPDPP---HPREGQKLLLHCEGRGNPVPQQYLWEK-
                                                                                                                                                                                                                                                                                                                                     SSGSKPAARLTWRKGDQELHGEPTRIQEDPNGKTFTVSSSVTFQVTREDDGASIVCSVNH
                                                                                                                                                                                                                                EGSVPPLKMTQESALIFPFLNKSDSGTYGCTATSNMGSYKAYYTLNVNDPS----PVPSSS
                                                                                                                                                                                                                                                                                                           PAVTG-NLOTORYLEVOYKPOVHIOMTYPLOGLTREGDALELTCEAIGKPOPVMVTWVRV 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   105;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AUG-1999) to the EMBL/GenBank/DDBJ databases
Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     432 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                         PRELIMINARY;
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 Chordata;
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                                                   Last sequence update)
Last annotation update)
                                                                             Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 469.5; DB 4;
Pred. No. 4.4e-31;
4; Mismatches 122;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Craniata; Vertebrata; I
Catarrhini; Hominidae;
 Craniata; Vertebrata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRT;
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 Euteleostomi;
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pfam; PF00047; 1g; 2.
SMART; SM00409; IG; 1.
SMART; SM00410; IG_11ke; 1.
SEQUENCE 549 AA; 60703 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
NCBI_TaxID=10090;
[1] ·
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL; AK011949; BAB27
MGD; MGI:1914402; 261
InterPro; IPR003599;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lyons P., Marchionni L., Mashima J., Mazzarelli J., Mombaerts P., Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N., Sasaaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F., Suzuki H., Toyo-oka K., Wang K.H., Weitz C., Whittaker C., Wilming L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=C57BL/6J; TISSUE=EMBRYO; MEDLINE=21085660; PubMed=11217851;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A. STRAIN=C57BL/6J; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nature 409:685-690(2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wynshaw-Boris
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YISDPPTTTTLQPTVQWHSSPADVQDIATEHKKLPFPLSTLATLKDDTIGTIIASVVGG
                                                                                   YVYDPP--TTIPP-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GPDSSIDGGNETVAAVCVSSTGKPVAQIDW-EGDLGEREFSTISFLNETATIVSQYELFP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           QKDTAVEG--EEIEVNCTAMASKPATTIRWFKGNTELKGKSEVEEWSDMYTVTSQLMLKV 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       QGRVLFKNYSLNDATITLHNIGFSDSGKYICKAVTFPLGNAQSSTTVTVLVEPTVSLIK-
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                                                                                                                                                                      DANPPPFKSVWSRLDGQWPDGLLASDNTLHFVHPLTVNYSGVYVCKVSNSLGQRSDQKVI
                                                                                                                                                                                                                                                          IGKPQPVMVTWVRVDDEMPQHAVLSGPNL-FINNLNKTDNGTYRCEASNIVGKAHSDYML
                                                                                                                                                                                                                                                                                                                                     TRFARGRRITCVVKHPALEKDIRYSFILDIQYAPEVSVTGYDGNWFVGRKG--VNLKCNA
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IPR003600; Ig_like.
IPR003006; Ig_MHC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2610301B19Rik.
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25.98;
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Pred. No. 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   32775CBEC7319B32 CRC64;
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                                                                                   -----PTTTTTTTTTTTLTILTITDSRAG
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.8e-17;
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Search completed: November Job time: 32.5 secs

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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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-MODEL-frame+_n2p_model -DEV-xlp
-Q-/cgn21/USPTO_spool/US9778187/runat_20112002_073631_26345/app_query.fasta_1.2318
-Q-/cgn21/USPTO_spool/US9778187/runat_20112002_073631_26345/app_query.fasta_1.2318
-DB-ISSUEd_Patents_AA -QFMT-fastan -SUFFIX-rai -MINMATCH-0.1 -LOOPCL-0
-LOOPEXT-0 -UNITS-bits -START-1 -END--1 -MATRIX-blosum62 -TRANS-buman40.cdi
-LIST-45 -DOCALIGN-200 -THR_SCORE-pct -THR_MS-100 -MIN-LIN-100-0 -ALIGN-15
-MODE-LOCAL -OUTFMT-pto -NORM-ext +HEAPSIZE-500 -MIN-LINE-0 -MAXLEN-2000000000
-USER-US09778187_eCGN_1_1_28_erunat_20112002_073631_26345 -NCPU-6 -ICPU-3
-USCR-US09778187_eCGN_1_1_28_erunat_20112002_073631_26345 -NCPU-6 -ICPU-3
-NORM-TIMEOUT-30 -THREADS-1 -XGAPPOF-10 -XGAPPEXT-0.5 -FGAPOP-6 -FGAPPEXT-7
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/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
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12,	Sequence 12, App	e 12,	e 33,	e 29,	e 16,	62,	21,	Sequence 19, App	e 15,		60,	2, 7	25,	e 17,	Sequence 45, App	49,	æ	4	43,	42,	e 47,	Ĺ	Sequence 3, Appl.	44,	48,	Sequence 2, Appl.	41,	Sequence 2, Appl	e 46,	e 2,	'n	Sequence 18, App
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### ALIGNMENTS

RESULT 1 US-08-659-984A-1

Sequence 1, Application US/08659984A Patent No. 5942400

GENERAL INFORMATION:

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APPLICANT: Anderson, John P.
APPLICANT: Jacobson-Croak, Kirsten L.
TITLE OF INVENTION: Assays for Detecting Beta-Secretase
TITLE OF INVENTION: Assays for Detecting Beta-Secretase
TITLE OF INVENTION: Inhibition
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Ctr., 8th Floor
CITY: San Francisco
STATE: California
COUNTR: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/659,984A
FILING DATE: 07-JUN-1996
CLASSIFICATION NUMBER: US 08/485,152
APPLICATION NUMBER: 15270-002810US
TELECOMMUNICATION NUMBER: 15270-002810US
TELECOMMUNICATION NUMBER: 15270-002810US
TELECOMMUNICATION NUMBER: 15270-002810US
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TELEFAX: 415-326-2422 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:

US-08-659-984A-1

MOLECULE TYPE:

protein

LENGTH: 421 amino acids TYPE: amino acid STRANDEDNESS: single TOPOLOGY: linear

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                                                                                                            GGGGATGCATTTGAGTTAACGTGTGAAGCCATCGGGAAGCCCCAGCCTGTGATGGTAACT
                                                                                                                                                       HisTyrThrProSerValLysIle-----IleProSerThrProPheProGlnGlu
                                                                                                                                                                          CAGTATAAACCGCAAGTGCATATCCAGATGACTTACCCTCTGCAAGGCCTAACCCCGGGAA
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    CTGTTCATCAATAACCTAAACAAAACAGATAACGGTACTTACCGCTGTGAGGCTTCCAAC
                                                              TGGGTGAGAGTCGATGAAAATG-----CCTCAACATGCCGTACTGTCTGGGCCAAAC
                                                                                         {\tt GlyGlnProLeuIleLeuThrCysGluSerLysGlyLysProLeuProGluProValLeu}
                                                                                                                                                                                                                  {\tt ArgValAspHisGluSerLeuAsnAlaThrProGlnValAlaMetGlnValLeuGluIlear}
                                                                                                                                                                                                                                             CAGGTGGAGCACCCTGCGGTCACTGGAAACCTGCAG---ACCCAGCGCTATCTAGAAGTG
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                               TrpThrLysAspGlyGlyGluLeuProAspProAspArgMetValValSerGlyArgGlu
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Query Match:
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APPLICANT: Chrysl
                                                                                                                                                                                                                           TELEFAX: 415-326-2422 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PM PC COMPATION
ODERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: 415-326-2400
                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/480,498
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Anderson, John P.
TITLE OF INVENTION: Beta-Secretase
NUMBER OF SEQUENCES: 21
                                                                                                                                  MOLECULE TYPE:
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                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS
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STATE: California
                                                                                                                                                                                                                                                      TELEPHONE:
                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                   NAME: Heslin, James M. REGISTRATION NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
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Result
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MODEL-frame+_n2p.model -DEV=xlp
-Q-/cgn2_1/USPTO_spool/US09778187/runat_20112002_073630_26329/app_query.fasta_1.2318
-Q-/cgn2_1/USPTO_spool/US09778187/runat_20112002_073630_26329/app_query.fasta_1.2318
-DB-PIR_73 -QFMT-fastan -SUPFIX-ppr -MIMMATCH-0.1 -LOOPCL-0 -LOOPEXT-0
-UNITS-bits -START-1 -END--1 -MATRIX-blosum62 -TRANS-human40.cdi -LIST-45
-DOCALIGN-200 -THR_SCORE-pct -THR_MAX=100 -THR_MIN-0 -ALIGN-15 -MODE-LOCAL
-OUTFMT-pto -NORM-ext -HEAPSIZE-500 -MINLEN-0 -MAXLEN-2000000000
-USER-US09778187_eCGN_1_1_113_@runat_20112002_073630_26329 -NCPU-6 -ICPU-3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES-0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS-1 -XGAPDP=10 -XGAPEXT-0.5 -FGAPDP=6 -FGAPEXT-7
                                                                                                                                                                                                                                                                                                                                                                                                                                               Database :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB seq
              Score
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2: pir2:*
3: pir3:*
4: pir4:*
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Match
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Listing first 45 summaries
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### ALIGNMENTS

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RESULT 1
T20992
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A;Reference number: Z19929
A;Accession: T24733
                            Query Match:
                                              Percent Similarity:
Best Local Similarity:
                                                                                                    Score:
                                                                                                                           Pred. No.:
                                                                                                                                               Alignment Scores:
                                                                                                                                                                                             A;Introns: 85/1; 120/1; 334/3; 370/1; 477/2; 606/3; 664/1; 935/3; 977/1; 1051/3; 1184; 2512/2; 2593/3; 2699/3; 2759/1; 2852/1; 2889/3; 2913/3; 2941/1; 2967/3; 2991/3; 3031; 4225/1; 4361/1; 4408/1; 4456/1; 4498/1; 4647/3; 4838/1; 4879/1; 4941/1; 5011/1; 50
                                                                                                                                                                                                                                                                     A; Map position: X
                                                                                                                                                                                                                                                                                               A; Gene: CESP:F15G9.4a
                                                                                                                                                                                                                                                                                                                                                A;Cross-references: EMBL:Z47070; PA;Experimental source: clone T09B9
                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: DNA
A; Residues: 1-5175 <WI2>
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A;Residues: 1-5175 <WIL>
A;Cross-references: EMBL:Z47068; PIDN:CAA87335.1; GSPDB:GN00028; CESP:F15G9.48
A;Experimental source: clone F15G9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: DNA
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A; Accession: T20992
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  4.16e-12
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                                                      {\tt IleThrValAsnSerGlyAsnProSerThrLeuHisCysProAlaLysGlySerProSer}
                                                                                                                                                                         AspPheLysValSerMetLeuValAlaProSerPheAspGluProAsnIleValArgArg
                                                                                                                                                                                                                                   GACTATATGCTGTATGTATACGATCCCCCCACAACTATCCCTCCT----
                                                                                                                                                                                                                                                                                            GluThrAspSerSerLysTyrThrCysIleAlaThrAsnGluAlaGlyThrAspLysArg
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GluGlyAspAlaGlyLysTyrThrCysGluAlaAspAsnSerAla-----GlySerVal 2372
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ACCATCCTTACCATCATCACAGATTCTCGAGCAGGTGAA
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                                                                                                                   -GlnAsnGluArgValValIleSerCysPro
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A; Notices: Fig. DNA A; Molecule type: DNA A; Residues: 1-5198 <WIL> A; Residues: 1-5198 <WIL> A; Cross-references: EMBL: Z47068; PIDN: CAA87336.1; GSPDB: GN00028; CESP: F15G9.4b
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: DNA
A; Residues: 1-5198 <WIZ>
A; Cross-references: EMBL: 247070; p.
A; Experimental source: clone T0989
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A; Residues: 1-5198 < VOG>
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A;Description: Hemicentin is required for hemidesmosome mediated
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hemicentin precursor - Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Date: 11-Jan-2000 #sequence_revision 11-Jan-2000
C;Accession: T43290; T20993; T24734
                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-778-187B-3_COPY_62_1069 (1-1008) x T43290 (1-5198)
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TTGATGATCGATATCCAGAAAGAC - - - ACGGCAGTTGAAGGGGAAGATTGAAGTCAAC
                                    As n Asp \verb|AlaGlyGluAsnSerPheAspPheLysAlaThrValLeuValProProThrIle| \\
                                                                                                                                                           AAAGTGTCACTGACGAATGTCTCAATCTCGGATGAAGGGAGATACTTCTGCCAGCTCTAC
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2967/3; 2991/3; 303
; 4964/1; 5034/1; 51
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-Q-/cgn2_l/USPTO_spool/US99778187/runat_20112002_073629_26309/app_query.fasta_1.2318
-Q-/cgn2_l/USPTO_spool/US997781887/runat_20112002_073629_26309/app_query.fasta_1.2318
-DB-SwissProt_40 -QFMT-fastan -SUFFIX=rsp -MINMATCH=0.1 -LOOPELT=0
-UNITS-bits -START=1 -END--1 -MATRIX=biosum62 -TRANS=human40.cd1 -LIST-45
-DOCALIGN-200 -THR_SCORE-pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE-LOCAL
-OUTFMT-pto -NORM-ext -HEARSIZE-500 -MINLEN0 -MAXLEN-200000000
-USER-US09778187_eCGN_1_145_erunat_20112002_073629_26309 -NOPU-6 -ICPU-3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-MARL_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPOEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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060469 homo sapien	O60500 homo sapien		Q08180 drosophila	w	homo s	P42292 gallus gall	rattus			P70211 mus musculu	homo s	gall	P13591 homo sapien		P13590 gallus gall	Q05793 mus musculu	mus mus	rattus	gallus	rattus	Q90610 gallus gall	mus m	homo	P06731 homo sapien	P35329 mus musculu	P97798 mus musculu

### ALIGNMENTS

RESULT 1

NCAL\_XENLA
ID NCAL\_XENLA
AC P16170;
DT 01-APR-1990 (Rel. 14, Created)
DT 01-APR-1990 (Rel. 14, Last sequence update)
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DT 15-JUN-2002 (Rel. 41, Last sequence update)
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RN SEQUENCE FROM N.A.
RX MEDLINE-90098071; PubMed=2481269;
RN SEQUENCE FROM N.A.
RX Krieg P. A., Sakayuchi D. S., Kintner C.R.;
RX Krieg P. A., Sakayuchi D. S., Kintner C.R.;
RX Krieg P. A., Sakayuchi D. S., Kintner C.R.;
RX Krieg P. A., Sakayuchi D. S., Kintner C.R.;
RX Krieg P. A., Sakayuchi D. S., Kintner C.R.;
RX MEDLINE-90098071; PubMed=2481269;
RX MEDLINE-90098071; PubMed This SWISS-PROT entry is copyright. It is produced through a collaboration between the SWISS-INSTITUTE of Bioinformatics and the EMBL outstation -Xenopus laevis (African clawed frog). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae; SUBCELLULAR LOCATION: Type I membrane protein.
ALTERNATIVE PRODUCTS: 2 isoforms; N-CAM 180; Shown here) and
N-CAM 180; are produced by alternative splicing.
TISSUE SPECIFICITY: EXPRESSED IN NEURON AND IN PRESUMPTIVE NEURAL EARLY NEURAL DEVELOPMENT.
SIMILARITY: BELONGS TO THE IMMUNOGLOBULIN SUPERFAMILY.
SIMILARITY: CONTAINS 5 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAIN
SIMILARITY: CONTAINS 2 FIBRONECTIN TYPE III-LIKE DOMAINS. FASCICULATION, ADHESION MOLECULE INVOLVED OF There are no restrictions LD-NCAM IS THE MAJOR IN THE EMBRYO DURING of a large l cell adhesion precursor (N-CAM DOMAINS n 09

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GGGAGATACTTCTGCCAGCTCTACACGGACCCCCCACAGGAGAGTTACACCACCATCACA
                 _IleSerPheAsnGluAspGlnSerGluMetThrIleHisHisValGluLysAspAspGlu
                                    CTGAATTTTTCTAGCAGTGAACTCAAAGTGTCACTGACGAATGTCTCAATCTCGGATGAA
                                                       AspProGluIleSerTrpLeuLysLysGlyGluProIleGluAspGlyGlu---GluLys
                                                                           AACAGGCAGACCATTTACTTCAGGGACTTCAGGCCTTTGAAGGACAGCAGGTTTCAGCTG
                                                                                            \tt AsnAlaThrAlaAsnMetAlaGluSerValValLeuSerCysAspAlaAspGlyPhePro
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                                                                                                                                 {\tt LysAspIleGlnValIleValAsnValProProThrIleGlnAlaArgGlnLeuArgValue} \\
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IPR003006; I9_MHC.
IPR003598; I9_c2.
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CYTOPLASMIC (POTENTIAL).

IG-LIKE C2-TYPE DOMAIN 1.

IG-LIKE C2-TYPE DOMAIN 2.

IG-LIKE C2-TYPE DOMAIN 3.

IG-LIKE C2-TYPE DOMAIN 4.

IG-LIKE C2-TYPE DOMAIN 4.
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FIBRONECTIN TYPE-III 2.
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HEPARIN-BINDING (POTENT
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62738B55B03F3E83
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Matches:
Conservative:
Mismatches:
Indels:
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Q9GL76;
16-OCT-2001 (Rel. 40, Created)
16-CCT-2001 (Rel. 40, Last sequence upda
15-JUN-2002 (Rel. 41, Last annotation up
Poliovirus receptor related protein 1 pu
mediator C) (HVeC) (Nectin 1).
PVRLI OR PRRI OR HVEC.
                                                                       SEQUENCE FROM N.A.
MEDLINE=21176378; PubMed=11277703;
Milne R.S.B., Connolly S.A., Krumm
Cohen G.H.;
                                                                                                                                                                                                               Sus scrofa (Pig).
Eukaryota; Metazoa; Chordata; Craniata; Ver:
Mammmalia; Eutheria; Cetartiodactyla; Suina;
"Porcine HveC, a member of the his a functional alphaherpesvirus Virology 281:315-328(2001).
                                                                                                                                                                                           NCBI_TaxID=9823;
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Sus.
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Command line parameters:

-MODEL-frame+ n2p.model -DEV-xlp
-Or/cgn2_1/USPTO_Spool/US09778187/runat_20112002_073633_26418/app_query.fasta_1.2318
-OP-Cgn2_1/USPTO_Spool/US09778187/runat_20112002_073633_26418/app_query.fasta_1.2318
-DB-Published_Applications_AA -OFMT-fastan -SUFFIX-rapb -MINMATCH-0.1
-LOOPEXT-0 -UNITS-bits -START=1 -END--1 -MATRIX-blosum62
-TRANS-human40.cdi -LIST-45 -DOCALIGN-200 -THR_SCORE-pct -THR_MAX=100
-THR_MIN-0 -ALIGN-15 -MODE-LOCAL -OUTFMT-pto -NORM-ext -HEAPSIZE-500 -MINLEN-0
-MAXLEN-200000000 -USER-US09778187 @CGN 1_1_15_@runat_20112002_073633_26418
-NCPU-6 -ICPU-3 -NO_XLPXY -NO_MMAP -LARGEOUERY -NGS_SCORES-0 -WAIT -LONGLOG
-DEV_TIMEOUT-120 -WARN_TIMEOUT-30 -THREADS-1 -XGAPOP-10 -XGAPEXT-0.5 -FGAPOP-6
-FGAPEXT-7 -YGAPOP-10 -YGAPEXT-0.5 -DELOP-6 -DELEXT-7
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Perfect score:
Sequence:
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                                                                                                                                                                                                                  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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1804
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97.8
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Listing first 45 summaries
                                                                                                                                                                                                                                                                                                : /cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
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: /cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep:*
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4, Appli
61, Appl
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#### ALIGNMENTS

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US-09-778-510-22

Sequence 22, Application US/09778510

Patent No. US20020164686A1

GENERAL INFORMATION:
APPLICANT: Baum, Peter
TITLE OF INVENTION:
APPLICANT APPLICATION NUMBER: US/09/778,510

CURRENT FILING DATE: 2001-02-07

PRIOR APPLICATION NUMBER: PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1999-08-05

PRIOR APPLICATION NUMBER: 60/095,663

PRIOR FILING DATE: 198-08-07

VOUNBER OF SEQ ID NOS: 22

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 22

LENGTH: 423

TYPE: PRT
ORGANISH: Mus musculus
US-09-778-510-22

Alignment Scores: 1.31e-115

Pred. No.: 1765.00

Best Local Similarity: 100.00%

Mismatches: 0

Query Match: 0

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Mismatches: 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/778,1878
CURRENT FILING DATE: 2001-02-06
PRIOR APPLICATION NUMBER: PCT/US99/17905
PRIOR FILING DATE: 1999-08-05
PRIOR APPLICATION NUMBER: US 60/095,672
PRIOR FILING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Baum, Peter R.
APPLICANT: Fanslow III, Willi
TITLE OF INVENTION: MOLECULES
FILE REFERENCE: 2873-US
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ORGANISM: mus
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P2548PIC1
FILE REFERENCE: P2548PIC1
CURRENT APPLICATION UNMBER: U$/09/944,413
CURRENT FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: December: 09/866,028
PRIOR FILING DATE: December: 11, 1997
PRIOR APPLICATION NUMBER: 60/667,411
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December: 11, 1997
PRIOR APPLICATION NUMBER: 60/069335
PRIOR FILING DATE: December: 11, 1997
PRIOR APPLICATION NUMBER: 60/069,325
PRIOR FILING DATE: December: 11, 1997
PRIOR APPLICATION NUMBER: 60/069,298
PRIOR FILING DATE: December: 11, 1997
PRIOR APPLICATION NUMBER: 60/069,425
PRIOR FILING DATE: December: 12, 1997
PRIOR APPLICATION NUMBER: 60/069,696
PRIOR FILING DATE: December: 10, 1997
PRIOR APPLICATION NUMBER: 60/069,696
PRIOR FILING DATE: December: 16, 1997
PRIOR APPLICATION NUMBER: 60/069,696
PRIOR FILING DATE: December: 16, 1997
PRIOR APPLICATION NUMBER: 60/069,702
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             DR FILING DATE: December 11, 1997
DR APPLICATION NUMBER: 60/069, 425
DR FILING DATE: December 12, 1997
DR APPLICATION NUMBER: 60/069,696
DR FILING DATE: December 16, 1997
DR APPLICATION NUMBER: 60/069,694
DR FILING DATE: December 16, 1997
DR APPLICATION NUMBER: 60/069,702
DR FILING DATE: December 16, 1997
DR APPLICATION NUMBER: 60/069,702
DR FILING DATE: December 16, 1997
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Wood, William
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Goddard, Audrey
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Percent Similarity:
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PRIOR APPLICATION NUMBER: 09/216,021
PRIOR APPLICATION NUMBER: 09/216,021
PRIOR PRILING DATE: December 16, 1998
PRIOR PELICATION NUMBER: 09/214,517
PRIOR PELICATION NUMBER: 09/254,311
PRIOR FILING DATE: March 3, 1999
PRIOR PRIOR DATE: Warch 3, 1999
PRIOR PELICATION NUMBER: PCT/US99/12252
PRIOR PELICATION NUMBER: PCT/US99/21090
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR PELICATION NUMBER: PCT/US99/2409
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TYPE: PRT
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OR APPLICATION NUMBER: PCT/USO0/04414
OR FILING DATE: February 22, 2000
OR APPLICATION NUMBER: PCT/USO0/05841
OR FILING DATE: March 2, 2000
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OR FILING DATE: March 30, 2000
OR APPLICATION NUMBER: PCT/USO0/14042
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FILING DATE: December 22, 1998
APPLICATION NUMBER: 60/146,222
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Baker,

Botstein, David

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US-09-944-403-61
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; Patent No. US20020165143A1
; GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: 05/09/944,403
CURRENT FILING DATE: 2001-05-26
PRIOR APPLICATION NUMBER: 05/066,028
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,335
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,278
PRIOR FILING DATE: December 12, 1997
PRIOR APPLICATION NUMBER: 60/069,425
PRIOR FILING DATE: December 12, 1997
PRIOR APPLICATION NUMBER: 60/069,696
PRIOR FILING DATE: December 16, 1997
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PRIOR APPLICATION NUMBER: 60/069,870
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PRIOR APPLICATION NUMBER: 60/074,086
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PRIOR APPLICATION NUMBER: 60/071,286
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PRIOR APPLICATION NUMBER: 60/0112,850
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PRIOR FILING DATE: DECEMBER: 60/146,222
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TITLE OF INVENTION: SECRETED
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PRIOR FILING DATE: February 28, 2001
NUMBER OF SEQ ID NOS: 120
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OR APPLICATION NUMBER: PCT/US00/05841
OR FILING DATE: MARCh 2, 2000
OR APPLICATION NUMBER: PCT/US00/08439
OR FILING DATE: MARCh 30, 2000
OR APPLICATION NUMBER: PCT/US00/14042
OR FILING DATE: May 22, 2000
OR APPLICATION NUMBER: PCT/US00/20710
OR FILING DATE: May 28, 2000
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FILING DATE: December 16, 1999
APPLICATION NUMBER: PCT/US00/03565
FILING DATE: February 11, 2000
APPLICATION NUMBER: PCT/US00/04414
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FILING DATE: No. US20020165143Alember
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APPLICANT: ROY, MAIGATEL
APPLICANT: Tumas, Daniel
APPLICANT: Tumas, Daniel
APPLICANT: WOOD, William
TITLE OF INVENTION: SECRETED AND TRANSME!
TITLE OF INVENTION: ACIDS ENCODING THE:
TITLE OF INVENTION: ACIDS ENCODING
CURRENT FILING DATE: DOCUMBER: 60,069,334
PRIOR APPLICATION NUMBER: 60,069,335
PRIOR APPLICATION NUMBER: 60,069,278
PRIOR APPLICATION NUMBER: 60,069,696
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Goddard, Audrey
Goddowski, Paul
Grimaldi, Christopher
Gurney, Austin
Hillan, Kenneth
Kljavin, Ivar
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OR APPLICATION NUMBER: 60/074.00
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APPLICATION NUMBER: PCT/US99/21090
FTI.ING DATE: September 15, 1999
FTI.ING DATE: September 16, 1999
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FILING DATE: No. US20020188715Alember
APPLICATION NUMBER: PCT/US99/2813
FILING DATE: No. US20020168715Alember
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FILING DATE: July 28, 1999
APPLICATION NUMBER: PCT/US98/19330
FILING DATE: September 16, 1998
APPLICATION NUMBER: PCT/US98/25108
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Best Local Similarity:
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TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P.2548P1C1
CURRENT APPLICATION NUMBER: US/09/866,028
CURRENT FILING DATE: 2001-05-25
Prior application data removed - consult PALM C
NUMBER OF SEQ ID NOS: 120
SEQ ID NO 61
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Gerritsen, Mary
Goddard, Audrey
Godowski, Paul
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Tumas, Daniel
Wood, William
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE PO TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P.2548P1C1
CURRENT APPLICATION NUMBER: US/09/944,449
CURRENT ELING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 09/866,028
PRIOR FILLING DATE: 2001-05-25
PRIOR FILLING DATE: December 3, 1997
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILLING DATE: December 11, 1997
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US-09-944-449-61
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Grimaldi, Christopher
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Goddard, Audrey
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E: December 16, 1998
N NUMBER: 60/113,296
E: December 22, 1998
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N NUMBER: 60/146,222
E: July 28, 1999
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N NUMBER: 60/074,086
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E: December 16, 1998
N NUMBER: 09/218,517
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NUMBER: 60/069,702
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NUMBER: PCT/US01/06520
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: June 22, 1999
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: September 15, 1999
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NUMBER: 60/075,945
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NUMBER: 60/069,694
: December 16, 1997
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NUMBER: 60/070,440
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CURRENT APPLICATION NUMBER: US/09/944,457
CURRENT FILING DATE: 2001-09-26
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R FILLING DATE: December 17, 1997
DR APPLICATION NUMBER: 60/069,873
DR FILLING DATE: December 17, 1997
DR APPLICATION NUMBER: 60/060,017
DR FILLING DATE: December 18, 1997
DR APPLICATION NUMBER: 60/070,440
DR FILLING DATE: Dacember 18, 1997
DR APPLICATION NUMBER: 60/070,440
DR FILLING DATE: January 5, 1998
DR APPLICATION NUMBER: 60/074,086
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R FILING DATE: December 16, 1997
DR APPLICATION NUMBER: 60/069,694
PR FILING DATE: December 16, 1997
DR APPLICATION NUMBER: 60/069,702
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                                                                              FILING DATE: February 25, 1 APPLICATION NUMBER: 60/112,
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FILING DATE: December 16, 1998
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FILING DATE: December 22, 1998
RAPPLICATION NUMBER: 09/254,311
FILING DATE: March 3, 1999
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FILING DATE: February 11, 2000
APPLICATION NUMBER: PCT/US00/04414
FILING DATE: February 22, 2000
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APPLICATION NUMBER: PCT/US99/28313
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APPLICATION NUMBER: PCT/US99/28301
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APPLICATION NUMBER: PCT/US00/14042
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INVENTION:
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                                       Wood, William
                                                       Roy,Margaret
Tumas,Daniel
                                                                                                            Hillan, Kenneth
Kljavin, Ivar
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                                                                                            Napier, Mary
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                                                                                                                                                                                         Godowski, Paul
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PRIOR APPLICATION NUMBER: 09/
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OR APPLICATION NUMBER: 60/069,702
OR FILING DATE: December 17, 1997
OR APPLICATION NUMBER: 60/069,870
OR FILING DATE: December 17, 1997
OR APPLICATION NUMBER: 60/068,017
OR APPLICATION NUMBER: 60/068,017
OR APPLICATION NUMBER: 60/068,017
OR FILING DATE: December 18, 1997
OR APPLICATION NUMBER: 60/070,40
OR FILING DATE: January 5, 1998
OR APPLICATION NUMBER: 60/074,092
OR FILING DATE: February 9, 1998
OR APPLICATION NUMBER: 60/075,945
OR FILING DATE: February 25, 1998
OR APPLICATION NUMBER: 60/075,945
OR FILING DATE: February 25, 1998
OR APPLICATION NUMBER: 60/112,850
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SR FILING DATE: December 16, 1998
OR APPLICATION WIMBER: 09/218,517
OR FILING DATE: December 22, 1998
OR APPLICATION WIMBER: 09/254,311
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                                                                                                                                                                                     APPLICATION NUMBER: PCT/US99/28313
FILING DATE: No. US20020115145A1eml
APPLICATION NUMBER: PCT/US99/28301
FILING DATE: December1, 1999
APPLICATION NUMBER: PCT/US99/30095
FILING DATE: December 16, 1999
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FILING DATE: September 15, 1980
APPLICATION NUMBER: PCT/US99/28409
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APPLICATION NUMBER: PCT/US99/12252
FILING DATE: June 22, 1999
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DATE: March 2, 2000
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Percent Similarity:
Best Local Similarity:
Query Match:
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PRIOR FILING DATE: July 28, 2000
PRIOR FILING DATE: PCT/US00/32678
PRIOR FILING DATE: December 1, 2000
PRIOR APPLICATION NUMBER: PCT/US01/06520
PRIOR FILING DATE: February 28, 2001
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CURRENT FILING DATE: US/09/945,587
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 09/866,028
PRIOR FILING DATE: 2001-05-25
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,325
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,278
PRIOR FILING DATE: December 12, 1997
PRIOR APPLICATION NUMBER: 60/069,425
PRIOR FILING DATE: December 16, 1997
PRIOR APPLICATION NUMBER: 60/069,696
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IOR APPLICATION NUMBER: 09/216,021
IOR APPLICATION NUMBER: 09/216,517
IOR FILING DATE: December 16, 1998
IOR APPLICATION NUMBER: 09/254,311
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IOR APPLICATION NUMBER: PCT/US99/1225
IOR APPLICATION NUMBER: PCT/US99/1225
IOR APPLICATION NUMBER: PCT/US99/21090
IOR FILING DATE: March 3, 1999
IOR APPLICATION NUMBER: PCT/US99/21090
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Eaton, Dan
Perrara, Napoleone
Pelivaroff, Ellen
Gerritsen, Mary
Goddard, Audrey
Godowski, Paul
Grimaldi, Christophe
                                                           Baker,
                                                   Botstein, David
                                                           Kevin
,Christopher
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 09/866,028
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
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CURRENT FILING DATE: 2001-09-26
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OR FILING DATE: December 18, 1997
OR APPLICATION NUMBER: 60/070,440
OR FILING DATE: January 5, 1998
OR APPLICATION NUMBER: 60/074,086
OR FILING DATE: FEBRUARY 9, 1998
OR APPLICATION NUMBER: 60/075,945
OR FILING DATE: FEBRUARY 9, 1998
OR APPLICATION NUMBER: 60/075,945
OR FILING DATE: FEBRUARY 25, 1998
OR APPLICATION NUMBER: 60/112,850
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 60/13,296
OR FILING DATE: December 22, 1998
OR APPLICATION NUMBER: 60/146,222
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 60/146,222
OR APPLICATION NUMBER: 60/146,222
OR APPLICATION NUMBER: 60/146,222
OR FILING DATE: September 16, 1998
OR APPLICATION NUMBER: FCT/US98/19330
OR APPLICATION NUMBER: FCT/US98/19330
OR APPLICATION NUMBER: FCT/US98/19330
OR APPLICATION NUMBER: FCT/US98/19330
                                                                                                                                                                                                                                                 OR APPLICATION NUMBER: 09/216,021
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 09/218,517
OR FILING DATE: December 22, 1998
OR APPLICATION NUMBER: 09/254,311
OR FILING DATE: March 3, 1999
OR APPLICATION NUMBER: PCT/US99/12252
OR FILING DATE: MIMBER: PCT/US99/21090
OR FILING DATE: September 15, 1999
OR APPLICATION NUMBER: PCT/US99/21090
OR FILING DATE: September 15, 1999
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DR APPLICATION NUMBER: 60/069,702
DR FILING DATE: December 16, 1997
DR APPLICATION NUMBER: 60/069,870
DR FILING DATE: December 17, 1997
DR APPLICATION NUMBER: 60/069,873
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APPLICATION NUMBER: 60/069,696
FILING DATE: December 16, 1997
APPLICATION NUMBER: 60/069,694
APPLICATION NUMBER: 60/069,694
                                                                                                     APPLICATION NUMBER: PCT/US99/28409
ETILING DATE: No. US20020132768Alember
APPLICATION NUMBER: PCT/US99/28313
FILING DATE: No. US20020132768Alember
APPLICATION NUMBER: PCT/US99/28301
ETILING DATE: No. US20020132768Alember
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APPLICATION NUMBER: 60/069,278
APPLICATION NUMBER: 61/11,1997
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APPLICATION NUMBER: 60/069,425
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Hillan, Kenneth
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Query Match:
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PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: February 22, 2000
PRIOR FILING DATE: February 22, 2000
PRIOR APPLICATION NUMBER: PCT/US00/05841
PRIOR FILING DATE: March 2, 2000
PRIOR FILING DATE: March 30, 2000
PRIOR PRIOR APPLICATION NUMBER: PCT/US00/14042
PRIOR APPLICATION NUMBER: PCT/US00/20710
PRIOR FILING DATE: May 22, 2000
PRIOR APPLICATION NUMBER: PCT/US00/20710
PRIOR APPLICATION NUMBER: PCT/US00/32678
PRIOR APPLICATION NUMBER: PCT/US00/32678
PRIOR APPLICATION NUMBER: PCT/US00/32678
PRIOR APPLICATION NUMBER: PCT/US00/32678
PRIOR FILING DATE: December 1, 2000
PRIOR FILING DATE: December 1, 2000
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PRIOR APPLICATION NUMBER: 09/866,028
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
PRIOR FILING DATE: December 11, 1997
PRIOR FILING DATE: December 11, 1997
PRIOR FILING DATE: December 11, 1997
PRIOR PRIOR PRICATION NUMBER: 60/069,334
PRIOR PRICATION NUMBER: 60/069,335
PRIOR APPLICATION NUMBER: 60/069,278
PRIOR APPLICATION NUMBER: 60/069,278
PRIOR APPLICATION NUMBER: 60/069,425
PRIOR APPLICATION NUMBER: 60/069,425
PRIOR APPLICATION NUMBER: 60/069,694
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CURRENT APPLICATION NUMBER: US/09/944,396
CURRENT FILING DATE: 2001-09-26
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                                                                                                                                       Query Match:
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OR FILING DATE: December: 09/218,51,
IOR APPLICATION NUMBER: 09/218,51,
IOR FILING DATE: December: 22, 1998
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The string Date: March 3, 1999
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DR APPLICATION NUMBER: 60/074,086
DR FILING DATE: February 9, 1998
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FILING DATE: February 25, 15.
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FILING DATE: December 16, 1998
APPLICATION NUMBER: 09/218,517
FILING DATE: December 22, 1998
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NUMBER: PCT/US00/32678
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RESULT 13
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; Patent No. US20020133675A1
; GENERAL INFORMATION:
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APPLICANT: Baker, Kevin
APPLICANT: Eaton, David
APPLICANT: Eaton, David
APPLICANT: Eaton, Dan
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APPLICANT: Eaton, Dan
APPLICANT: Ferrara, Mappleone
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APPLICANT: Gerritsen, Mary
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APPLICANT: Hillan, Kenneth
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PRIOR FILING DATE: May 22, 2000
PRIOR APPLICATION NUMBER: PCT/US00/20710
PRIOR FILING DATE: July 28, 2000
PRIOR APPLICATION NUMBER: PCT/US00/32678
PRIOR APPLICATION NUMBER: PCT/US01/06520
PRIOR APPLICATION NUMBER: PCT/US01/06520
PRIOR APPLICATION NUMBER: PCT/US01/06520
PRIOR FILING DATE: February 28, 2001
NUMBER OF SEQ ID NOS: 120
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OR APPLICATION NUMBER: PCT/US99/28301
OR FILING DATE: December1, 1999
OR APPLICATION NUMBER: PCT/US99/30095
OR FILING DATE: December 16, 1999
OR APPLICATION NUMBER: PCT/US09/03565
OR FILING DATE: February 11, 2000
IOR APPLICATION NUMBER: PCT/US00/04414
OR FILING DATE: February 22, 2000
IOR APPLICATION NUMBER: PCT/US00/05841
OR FILING DATE: MECHUARY 22, 2000
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FILING DATE: March 30, 2000
APPLICATION NUMBER: PCT/US00/08439
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                      TGGTTCAAAGGGAACAAGGAACTCAAAGGCAAATCAGAGGTGGAGGAGTGGTCGGACATG
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APPLICATION WOOD, WILLIAM
APPLICATION: SECRETED AND TRANSMEMBRANE POL
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POL
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: p.2548P1C1
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: US/09/44,432
CURRENT FILING DATE: 2001-05-25
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/667,411
PRIOR FILING DATE: December 3, 1997
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069335
PRIOR APPLICATION NUMBER: 60/069,278
PRIOR APPLICATION NUMBER: 60/069,278
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,278
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,278
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Goddard, Audrey
Goddwski, Paul
Grimaldi, Christopher
Gurney, Austin
Hillan, Kenneth
Kljavin, Ivar
Napler, Mary
Roy, Margaret
Tumas, Daniel
Wood, William
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(OR APPLICATION NUMBER: PCT/US99/12252

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OR APPLICATION NUMBER: 60/069,017
OR APPLICATION NUMBER: 60/069,017
OR APPLICATION NUMBER: 60/070,440
OR FILING DATE: December 18, 1997
OR APPLICATION NUMBER: 60/070,440
OR FILING DATE: January 5, 1998
OR APPLICATION NUMBER: 60/074,086
OR FILING DATE: February 9, 1998
OR APPLICATION NUMBER: 60/074,092
OR FILING DATE: February 9, 1998
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PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
PRIOR APPLICATION NUMBER: 60/069,334
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OR APPLICATION NUMBER: 60/075,945

OR FILING DATE: February 25, 1998

OR APPLICATION NUMBER: 60/112,850

OR FILING DATE: December 16, 1998

OR APPLICATION NUMBER: 60/113,296
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APPLICATION NUMBER: 60/069,694
FILING DATE: December 16, 1997
APPLICATION NUMBER: 60/069,702
FILING DATE: December 16, 1997
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APPLICATION NUMBER: PCT/US98/25108
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APPLICATION NUMBER: 60/074,08
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APPLICATION NUMBER: 60/069,278
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NUMBER: PCT/US98/19330
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Command line parameters:

-MODEL-frame+_n2p.model -DEV-xlp

-MODEL-frame+_n2p.model -DEV-xlp

-O-/cgn2_1/USPTO_spool/US9778187/runat_20112002_073629_26303/app_query.fasta_1.2318

-DB-A_Geneseq_101002 -OFMT-fastan -SUFFIX-rag -MINNATCH-0.1 -LOOPCL-0

-LOOPEXT-0 -UNITS-bits -START-1 -END--1 -MATRIX-blosum62 -TRANS-human40.cdi
-LIST-45 -DOCALIGN-200 -THR_SCORE-pct -THR_MAX-100 -THR_MIN-0 -ALIGN-15

-MODE-LOCAL -OUTFMT-pto -NORM-ext -HEAPSIZE-500 -MINLEN-0 -MAXCHEN-200000000

-USER-US09778187_eCGN_1_1_71_erunat_20112002_073629_26303 -NCPU-6 -ICPU-3

-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES-0 -WAIT -LONGLOG -DEV_TIMEOUT-120

-WARN_TIMEOUT-30 -THREADS-1 -NORM-PC -N
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## SUMMARIES

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## ALIGNMENTS

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DT 31-MAY-2000 (first entry)
XX

Expended derived dendritic cell adhesion molecule.
XX

Expended derived dendritic cell adhesion molecule; LDCAM; mouse; B7-1;
XX

Expended derived dendritic cell adhesion molecule; LDCAM; mouse; B7-1;
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Expended derived dendritic cell adhesion molecule; LDCAM; mouse; B7-1;
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                                                                                                                                                                                                                                                                                       cell adhesion molecule, LDCAM. It is found on lymphoid derived dendritic cells and displays homology to adhesion molecules, B7-1 and cytoplasmic region of B7-L1. Mouse LDCAM is found on whole embryo, testes, triple negative cells murine splenic and lymph node CD8+, S49.1 and dendritic cells. LDCAM polypeptides interacts with T cell surface molecules to alter signalling and inhibits T cell proliferation, bind to themselves and B7L-1, an LDCAM binding protein and increases natural killer (NK) cell populations. It may be used to measure the biological activity and as quality control reagents of LDCAM binding proteins. LDCAM may be used for treating disorders associated with malfunctioning of impure the protein of the protein and increase in the protein and increase in the protein and increase in the protein and increase the biological activity and as quality control reagents of LDCAM binding proteins.
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17-JUL-2000;
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15-SEP-2000;
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                                                                                                                                                                                                                                                                                                                polypeptide for treatment s and research use -
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2000US-0617746.
2000US-0631451.
2000US-0663B70.
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TrpPheLysGlyAsnThrGluLeuLysGlyLysSerGluValGluGluTrpSerAspMet
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98US-0075945.
97US-0067411.
97US-0069278.
97US-0069334.
                                       98WO-US25108
                                                                                                   transmembrane
                                                                                                                                                                      Protein;
                                                                                                          tumour necrosis factor family; TNF;
                                                                                                                         sequence
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05-JAN-1998
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                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes nucleic acids encoding PRO secreted and transmembrane proteins used therapeutically. The PRO proteins have cytostatic, anti-inflammatory, anti-proliferative and immunosuppressive activity. The proteins and polynucleotides can be used in therapy,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
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                                                                                                                                                                                                                                                                                                                                                                                                            complement
                                                                                                                                                                                                                                                                                                                                                                                                                identification of homologues, raising antibodies and design of probes and primers. They can be used in a range of diseases related to proteins that they have homology with, e.g. a PRO protein having homology to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Baker
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ACAGTCCTGGTTCCTCCACGTAACTTGATGATCGATATCCAGAAAGACACACGGCAGTTGAA
                                                                                          GAAGGGAGATACTTCTGCCAGCTCTACACGGACCCCCCACAGGAGAGTTACACCACCATC
                                                                                                                                  CTGCTGAATTTTTCTAGCAGTGAACTCAAAGTGTCACTGACGAATGTCTCAATCTCGGAT
                                                                                                                                                             {\tt ProAsnArgGlnThrIleTyrPheArgAspPheArgProLeuLysAspSerArgPheGln}
                                                                                                                                                                         CCCAACAGGCAGACCATTTACTTCAGGGACTTCAGGCCTTTGAAGGACAGCAGCAGTTTCAG
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DB; AAX80055.
                                         ThrValLeuValProProArgAsnLeuMetIleAspIleGlnLysAspThrAlaValGlu
                                                                                 GluGlyArgTyrPheCysGlnLeuTyrThrAspProProGlnGluSerTyrThrThrIle
                                                                                                                      {\tt LeuLeuAsnPheSerSerGluLeuLysValSerLeuThrAsnValSerIleSerAsp}
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97US-0069425.
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97US-0069696.
97US-0069702.
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97US-0068017.
98US-00704440.
98US-00744086.
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163..167
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9..15
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233..2
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65..69
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Gerritsen M
Hillan KJ,
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16-DEC-1998;
22-DEC-1998;
                                                                                                                  New human nucleic acids encoding secreted and transmembrane polypeptides which are designated as PRO polypeptides are described The membrane-bound proteins have various industrial applications, including as pharmaceutical and diagnostic agents. The membrane-bound proteins can also be employed for screening of potential peptide or small molecule inhibitors of the relevant receptor/ligand interaction. Anti-PRO antibodies are useful for the affinity purification of PRO from recombinant cell culture or natural sources.
                                                                                                                                                                                             Claim 12;
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J, Kljavin
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les, designated as PRO polypeptides, useful as pharmaceutical
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             Sequences
                                                       Novel nucleic acids encoding PRO polypeptides, used presence of tumours, such as prostate and breast tun to screen for modulators of the compounds -
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J, Smith V,
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2000WO-US34956.
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2000WO-US05841
2000WS-187202P
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2000US-19335P
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The present invention relates to 12 secreted human proteins and the nucleotide sequences encoding them. The polynucleotide sequences given in AAA80606-A80623 encode the 12 secreted protein sequences given in AAB25576-B2593. The human secreted proteins have various activities dependent on the tissues in which they are expressed. Examples of the activities of the proteins include: immunosuppressant; anti-inflammatory antiarthritic; antirheumatic, dermatological; antiproliferative; antiarteriosclerotic; anticancer; vulnerary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Secreted protein; immunosuppressant; anti-inflammatory; antiarthritic; antirheumatic, dermatological; antiproliferative; antiarteriosclerotic; anticancer; vulnerary; antiviral; antibacterial; antifungal; immune disorder; Addison's disease; rheumatoid arthritis; dermatitis; multiple sclerosis; inflammatory disorder; inflammatory bowel disease; crohn's disease; nephritis; hyperproliferative disorder; cancer; cardiovascular disorder; coronary arteriosclerosis; myocarditis; cancer; melanoma, sconar beating.
                                                                                                                                                                                                                                    Isolated nucleic acid molecules encoding human secreted proteins a used for the prevention, amelioration and treatment of autoimmune, inflammatory, hyperproliferative and cardiovascular disorders, carwounds, and infectious diseases.
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Best Local Similarity:
Query Match:
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Percent Similarity: Best Local Similarity: Query Match:

Conservative: Mismatches: Indels:

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Gaps:

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Alignment
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07-DEC-1998;
08-MAR-1999;
                                                                                                                                                                                                                                                                        akathesia, Alzheimer's disease, multiple sclerosis and epilepsy. Polynucleotides encoding HSCRPs may be used as hybridisation probe diagnose these conditions. Anti-HCSRP antibodies may be used as antagonists, as a targeting or delivery mechanism for bringing pharmaceutical agents into contact with cells or tissues expressing the state of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence is a novel human cell surface receptor protein (HCSRP) designated HCSRP-8. The nucleotide sequence was identified in Incyte Clone 312256 from the cDNA library LUNGNOTO2, which was made from RNA isolated from lung tissue. A number of Incyte Clones were used to assemble the consensus sequence. BLAST analysis showed that the sequence is homologous to immuno-superfamily protein B12 3779242. HCSRP and its
                                                                                                                                                                                                                                                                                                                                                                                                           disorders include cancers such as leukaemia and melanoma, immune disorders such as rheumatoid arthritis, asthma and atherosclerosis, bacterial and parasitic infections and neuronal disorders such as
                                                                                                                                                                                      Sequence
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Key
Domain
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ovel molecules designated LDCAM cell function are capable of altering or modulating

Claim 7; Page 42-43; 44pp; English

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Cells and displays homology to adhesion molecules, B7-1 and cytoplasmic crayion of B7-L1. Human LDCAM. It is found on lymphoid derived dendritic cells and displays homology to adhesion molecules, B7-1 and cytoplasmic CC region of B7-L1. Human LDCAM is expressed in breast, retina, foetal CC liver, spleen and heart, lung, muscle, placenta, thyroid and lung CC carcinoma. LDCAM polypeptides interacts with T cell surface molecules CC to alter signalling and inhibits T cell proliferation, bind to CC themselves and B7L-1, an LDCAM binding protein and increases natural CC killer (NK) cell populations. It may be used to measure the biological CC cells in and as quality control reagents of LDCAM binding proteins. CC LDCAM may be used for treating disorders associated with malfunctioning CC cells, infectious diseases and for killing tumour cells. They are also CC useful for prevention or reducing the effect of organ and bone marrow CC useful for prevention and for modulating T cell immune responses. LDCAM CC colls any also be used as carriers for delivering agents attached CC to T cells or cells bearing B7L-1. present amino acid sequence is the human lymphoid derived dendritic

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                                                       ACTTGGGTGAGAGTCGATGATGAAATGCCTCAACATGCCGTACTGTCTGGGCCAAACCTG
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                                                                                The invention relates to a method for detecting cell proliferative collsorder associated with tumour suppressor lung cancer 1 (TSLC1) in a subject. The method comprising contacting a cell component of a proliferating cell with a reagent that detects level of the cell component in the proliferating cell and determining modification in the level of the cell component in proliferating cell as compared with a cell cell component in proliferating cell as compared with a cell thy cell, where modification indicates disorder associated with TSLC1. The method is useful for detecting a cell proliferative disorder (e.g. liver, lung or pancreatic cancer) associated with tumour suppressor lung cancer 1 (TSLC1) in a subject. The invention is useful in gene therapy and for treating a cell proliferative disorder such as lung cancer (human non-small cell lung cancer), liver cancer (hepatocellular carcinoma) or pancreatic cancer associated with modification of TSLC1 production, where a reagent which modulates (preferably, increases) TSLC1 level in the cells is compared.
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in TSLC1 level -
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                                                                                 TTCATCAATAACCTAAACAAAACAGATAACGGTACTTACCGCTGTGAGGCTTCCAACATA
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Best Local S Query Match:

US-09-778-187B-3\_COPY\_62\_1069 (1-1008)

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(1-364)

Percent Similarity: Best Local Similarity:

4e-142 1694.00 99.38% 99.38% 93.90% 21

Length:
Matches:
Conservative:
Mismatches:
Indels:

364 0 2 0

Score:

NO . .

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The present invention relates to 12 secreted human proteins and the CC nucleotide sequences encoding them. The polynucleotide sequences given in AAA80606-A80623 encode the 12 secreted protein sequences given in CC AAB2576-B2593. The human secreted proteins have various activities CC dependent on the tissues in which they are expressed. Examples of the CC activities of the proteins include: immunosuppressant; CC antivitial antibacterial; antiarteriosclerotic; anticancer; vulnerary; CC antivital; antibacterial; and antifungal activity. The proteins, CC entivital; antibacterial; and antisqual activity. The proteins, CC polypeptides, agonists and antagonists may be used to treat prevent CC and/or diagnose various disease, disorders and conditions examples of CC which include: immune disorders e.g. Addison's disease, rheumatoid CC arthritis, dermatitis, and multiple sclerosis; inflammatory disorders CC e.g. inflammatory bowel disease, Crohn's disease and nephritis; CC hyperproliferative disorders such as paraproteinaemlas and purpura; CC cardiovascular disorders e.g. coronary arteriosclerosis and myocarditis; CC cancer e.g. melanoma and lymphoma. The proteins and polynucleotide sequences may also be used in wound healing and the treatment of consequences are represented in sequences AAA80616 and AAB25586. Sequences are represented in sequences AAA80616 and AAB25586. Sequences are represented in sequences and the treatment of the AAB25586.
Alignment
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                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Isolated nucleic acid molecules encoding human secreted proteins are used for the prevention, amelioration and treatment of autoimmune, inflammatory, hyperproliferative and cardiovascular disorders, cancer wounds, and infectious diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Greene
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                                         ThrAspSerArgAla
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US-09-778-187B-3\_COPY\_62\_1069 (1-1008)

Indels:
Gaps:
x AAB88427

(1-443)

Percent Similarity: Best Local Similarity:

4.54e-133 1592.50 92.88% 91.39% 88.28%

443 308 5 23 1

Length:
Matches:
Conservative:
Mismatches:

No.

Query

Match:

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CC which encode human secretory or membrane proteins represented by
CR AAB88317 - AAB8419. Included in the invention are primers
CR AAB88317 - AAB8419. Included in the invention are primers
CR AAF93917 - AAF94295 and AAF62232 - AAF62235 which are used to isolate the
CR CDNA sequences of the invention. The invention also includes methods for
CR the production of antibodies directed against the proteins, and CDNA
CR sequences, which can be used in vaccines. The polynucleotide sequences
CR can be used in gene therapy. The polynucleotide sequences and the
CR proteins they encode may be used in the prevention, treatment and
CR diagnosis of diseases associated with inappropriate secretory
CR sequences may also be used as DNA probes in diagnostic assays
CR (e.g. polymerase chain reactions (PCR)) to detect and quantitate the
CR presence of similar nucleic acid sequences in samples. They may also be
CR used to study the expression and function of secretory proteins/membrane
CR polypeptides and their role in metabolism. The polypeptides may be used
CR as antigens in the production of antibodies against them and in assays to
CR identify modulators (agonists and antagonists) of expression and
CR cactivity. The antibodies and antagonists may also be used as therapeutic
CR agents to down regulate expression and activity. The antibodies may also
CR be used as diagnostic agents for detecting the presence of the
CR polypeptides in samples (e.g. by enzyme linked immunosorbant assay
CR (ELISA). Examples of diseases which may be treated include rheumatoid
CR arthritis and diabetes.
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11-JAN-2000;
02-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention relates to nucleic acid sequences AAF93744 -
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rheumatoid arthritis; diabetes.
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as candidate target molecules in drug development -
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                                                                                                                                ATCACAGATTCTCGAGCAGGTGAAGAGGGGACCATTGGGGCAGTGGACCAC
                                                                                                                                                       ProAlaValHisGlyLeuThrGlnLeuProAsnSerAlaGluGluLeuAspSerGluAsp
                                                                                                                                                                            ValGlyLysAlaH1sSerAspTyrMetLeuTyrValTyrAspThrThrAlaThrThrGlu
                                                                                                                                                                                                                                                                                      {\tt ThrTrpValArgValAspAspGluMetProGlnHisAlaValLeuSerGlyProAsnLeu}
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DB:
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Best Local Similarity:
                                                        US-09-778-187B-3_COPY_62_1069 (1-1008)
                                                                                                                                              Score:
                                                                                                                                                           Pred.
                                                                                                                                                                       Alignment
                                                                                                                                                                                                                                      The present invention describes new human secreted proteins which were careful from adult placenta, adult trettana, foetal brain, foetal kidney, adult blood, adult thrain, adult tryroid, adult blood, adult neural careful kidney, adult blood, adult train, adult tlymph node cDNA libraries. The human careful from the polynucleotides encoding them, are predicted to have biological activities which would make them suitable for treating, preventing or ameliorating medical conditions in humans and animals. Suggested activities include nutritional activity, cytokine and cell proliferation/differentiation activity, immune stimulating (e.g. as vaccines) or suppressing activity, haematopoiesis regulating cactivity, tissue growth activity, activity, haematopoiesis regulating cactivity, tissue growth activity, activity, haemostatic and thrombolytic activity, receptor/ligand activity, anti-inflammatory activity, receptor/ligand activity, anti-inflammatory activity or activity. The polynucleotides are also stated to be useful for gene therapy. AAZ33316 to AAZ3373 encode human secreted proteins, given in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-JUN-1998;
22-JUL-1998;
31-JUL-1998;
10-AUG-1998;
11-AUG-1998;
06-MAY-1999;
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Merberg D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          haematopoiesis regulation; tissue growth; activin; inhibin; chemotactic; chemokinetic; haemostatic; thrombolytic; receptor; ligand; anti-inflammatory; cadherin; tumour invasion suppressor; tumour inhibition; gene therapy.
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differentiation; immune stimulating; vaccine; suppression;
haematopoiesis regulation; tissue growth; activin; inhibin;
chemotactic; chemokinetic; haemostatic; thrombolytic; receptor;
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sapiens
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DB; AAZ33346.
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98US-0095880.
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Conservative:
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Steininger RJ, 1
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RESULT 1:
AAB25593
ID AAB2
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DT 21-N
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DE Prot
                               21-NOV-2000
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multiple sclerosis; inflammatory disorder; inflammatory bowel disease; Crohn's disease; nephritis; hyperproliferative disorder; cardiovascular disorder; coronary arteriosclerosis; myocarditis; cancer.
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anticancer; vulnerary; antiviral; antibacterial; antifungal;
immune disorder; Addison's disease; rheumatoid arthritis; dermatitis;
                                                                                                                                                                                                                                                                          Isolated nucleic acid molecules encoding human secreted proteins are used for the prevention, amelioration and treatment of autoimmune, inflammatory, hyperproliferative and cardiovascular disorders, cancer, wounds, and infectious diseases
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Claim 1; Page 685-686; 803pp; English.

anti-inflammatory; antiarthritic; antirheumatic, dermatological; antiproliferative; antiarteriosclerotic; anticancer; vulnerary; antiviral; antibacterial; and antifungal activity. The proteins, polypeptides, agonists and antiagonists may be used to treat prevent and/or diagnose various disease, disorders and conditions examples of which include: immune disorders e.g. Addison's disease, rheumatoid arthritis, dermattits, and multiple sclerosis; inflammatory disorders e.g. inflammatory bowel disease, Crohn's disease and nephritis; hyperproliferative disorders such as paraproteinaemias and purpura; and purpura scheme and s nucleotide sequences encoding them. The polynucleotide sequences given in AAA80606-A80623 encode the 12 secreted protein sequences given in AAB25576-B25593. The human secreted proteins have various activities dependent on the tissues in which they are expressed. Examples of the activities of the proteins include: immunosuppressant; cardiovascular disorders e.g. coronary arteriosclerosis and myocarditis, cancer e.g. melanoma and lymphoma. The proteins and polynucleotide sequences may also be used in wound healing and the treatment of AAA80677-A80682 infectious diseases. The human secreted protein gene #11 and protein sequences are represented in sequences AAA80616 and AAB25586. Sequences present invention relates represent genes to 12 secreted human related 6 secreted protein proteins

Sequence

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                                                                                                                                                                                    Human; cytokine; ceil proliferation; cell differentiation; gene therapy; vaccine; peptide therapy; stem cell growth factor; haematopoiesis; tissue growth factor; humunomodulatory; cancer; leukaemia; nervous system disorder; arthritis; inflammation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to polynucleotides (AAK51456-AAK53435) and the encoded polypeptides (AAM78323-AAM80302) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polynucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopolesis regulating activity tissue growth factor activity, immunomodulatory activity and activin/inhibin activity and may be useful in the diagnosis and/or treatment of cancer, leukaemia, nervous system disorders, arthritis and
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                                    ValGluLeuSerCysLeuValProArgSerArgProAlaAlaThrLeuArgTrpTyrArg
                                                                               ValAlaProGluAsnProValValGluVal---ArgGluGlnAlaValGluGlyGlyGlu
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, Wang D,
Yang Y,
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T, Goodrich R;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                               Arg Val Asp His Glu Ser Leu Asn Ala Thr Pro Gln Val Ala Met Gln Val Leu Glu Ileu G
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HisTyrThrProSerValLysIle--

-IleProSerThrProPheProGlnGlu

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( <u>)</u>	370	976	350	958	330	898	310	838	290	778	270	724	250	664
Search completed: November 20. 2002. 07:50:19	GlyGlnAsnGlyProAspHis 376	ACCATTGGGGCAGTG	ThrThrSerProThrThrSerAlaThrThrSerSerIleArgAspProAsnAlaLeuAla 369	ATCACA	LeuProThrThrIleIleProSerLeuThrThrAlaThrValThrThrThrValAlaIle 349	CCTCCTCCCACAACAACCACC	ThrileGlyGlnSerSerAlaGluTyrValLeuIleValHisAspValProAsnThrLeu 329	ATA	LeuAsnIleLeuPheLeuAsnLysThrAspAsnGlyThrTyrArgCysGluAlaThrAsn 309	CTGTTCATCAATAA	270 TrpThrLysAspGlyGlyGluLeuProAspProAspArgMetValValSerGlyArgGlu 289	STCGATGATGAAATGCCTCAACATGCC	GlyGlnProLeuTleLeuThrCysGluSerLysGlyLysProLeuProGluProValLeu 269	GGGGATGCATTTGAG

Search completed: November 20, 2002, 07:50:19 Job time : 59.25~secs

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Command line parameters:

-MODEL-framet_n2p.model -DEV-xlp
-O-/cgn2_1/USPTO_spool/US09778187/runat_20112002_073630_26319/app_query.fasta_1.2318
-O-/cgn2_1/USPTO_spool/US09778187/runat_20112002_073630_26319/app_query.fasta_1.2318
-DB-SPTREMBL_21 -OFMT-fastan -SUFFIX=rspt -MINMATCH-0.1 -LOOPCL-0 -LOOPEXT-0
-UNITS-bits -START-1 -END--1 -MATRIX=blosum62 -TRANS-human40.cd1 -LIST-45
-DOCALIGN-200 -THR_SCORE-pct -THR_MAN-100 -ALIGN-15 -MODE-LOCAL
-OUTPWT-pto -NORM-ext -HEAPSIZE-500 -MIXLEN-2000000000
-USER-US09778187_eCGN_11_204_@runat_20112002_073630_26319 -NCPU-6 -ICPU-3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES-0 -WAIT -LONGLOG -DEV_TIMEOUT-120
-WARN_TIMEOUT-30 -THREADS-1 -FGAPOP-10 -XGAPEXT-0.5 -FGAPOP-6 -FGAPEXT-7
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Q8R4L1;
Q1-JUN-2002 (TrEMBLrel. 21, Created)
Q1-JUN-2002 (TrEMBLrel. 21, Last sequence update)
Q1-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Tumor suprressor in lung cancer 1.
                                                                                           Fukami T., Maruyama T., Murakami Y.;

"Identification of murine orthologue of the TSLC1

Submitted (OCT-2001) to the EMBL/GenBank/DDBJ data

EMBL; AP434663; AALB6736.1; -.

SEQUENCE 445 AA; 48664 MW; C5D5A070DAF70E55 CF
                                                                                                                                                                                                                                                                  Mus musculus (Mouse).
Eukaryota; Metazoa; C
Mammalia; Eutheria; R
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Rodentia;
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Length:
Matches:
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RA175.
Mus musculus
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RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
RA Altawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
Ra Altawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
Ra Altawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
Ra Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
Ra Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,
Ra Kuchi P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
Ra Kuchi P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
Ra Kuchi P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
Ra Schrimi L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,
Ra Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
Ra Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
Ra Bluke J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
Ra Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M.,
Ra Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
Ra Lyons P., Marchionni L., Mashima J., Mazzarelli J., Mombaerts P.,
Ra Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,
Ra Sasaki H., Sato K., Wang K.H., Weitz C., Whittaker C., Wilming L.,
Ra Mynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
Ra Hayashizaki Y.,
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O9CRY3;
O1-JUN-2001 (TrEMBLrel. 17, Created)
O1-JUN-2001 (TrEMBLrel. 19, Last sequence update)
O1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
O1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
O1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
IGSF4 OR 310000110BRIK.
Mus musculus (Mouse).
Ekkaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
NCBI_TaxID-10090;
                 EMBL; AK01
MGD; MGI:1
InterPro;
                                  Häyashizaki Y.;
"Functional annotation of a
Nature 409:685-690(2001).
EMBL; AK013911; BAB29050.1;
MGD; MGI:1889272; Igsf4.
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InterPro; IPR003006; Ig_MH
Pfam; PF00047; Ig; 3.
SMART; SM00409; IG; 3.
SMART; SM00408; IGC2; 3.
SMART; SM00410; IG_1ike; 1
Immunoglobulin domain.
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InterPro; IPR003508; Ig_like.
InterPro; IPR003006; Ig_MHC.
Pfam; PF00047; ig; 3.
SMART; SM00409; IG; 3.
SMART; SM00408; IG:2; 3.
SMART; SM00410; IG_like; 2.
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"Cloning of a novel human
superfamily.";
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Q9BY67;
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SEQUENCE 442
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Mammalia; Eutheria;
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Metazoa; Primates; (
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                         SEQUENCE FROM N.A. STRAIN-C57BL/6J; TISSUE-HIPPOCAMPUS; MEDLINE-21085660; PubMed-11217851;
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  Shinagawa
  Α.,
  Shibata
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Query Match:
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RA Alzawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Salto T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Salto R.,
RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
RA Kuchi R., Watsuda T., Gissi C., King B., Kochiwa H.,
RA Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,
RA Kuchi P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuchi P., Lewis S., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
RA Lyons P., Marchionni L., Mashima J., Mazzarelli J., Mombaerts P.,
RA Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,
RA Suzuki H., Toyo-oka K., Wang K.H., Weitz C., Whittaker C., Wilming L.,
RA Hayashizaki Y.,
RA Hayashizaki Y.,
RENEWARD R., Kanga Y., Kawaji H., Kohtsuki S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nature 409:685-690(2001).

EMBL; AK013775; BAB28988.1; -.

MGD; MGI:1889272; Igsf4.

InterPro; IPR003599; Ig_C2.

InterPro; IPR003598; Ig_C2.

InterPro; IPR003500; Ig_like.

InterPro; IPR003006; Ig_MHC.

Pfam; PF00047; Ig; 3.

SMART; SM00409; IG; 3.

SMART; SM00408; IG_C2; 3.

SMART; SM00408; IG_C2; 3.

SMART; SM00409; IG_C2; 3.

SMART; SM00410; IG_Like; 1.

Immunoglobulin domain.

SEQUENCE 336 AA; 37157 MW;
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TrpPheLysGlyAsnLysGluLeuLysGlyLysSerGluValGluGluTrpSerAspMet
                                TGGTTCANAGGGAACAAGGAACTCAAAGGCAAATCAGAGGTGGAGGAGTGGTCGGACATG
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Best Local Similarity:
Query Match:
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SEQUENCE FROM N.A.

Zhou Y., Du G., Chen J., Yuan J., Qla
Zhou Y., Du G., Chen J., Tuan J., Qla
"Cloning of a novel cDNA encoding a m
Submitted (APR-1998) to the EMBL/GenB
Submitted (APR-1998) to the EMBL/GenB
EMBL, AF061260, AAC67243.1; -.

MGD: MGI:1889272; Igsf4.

InterPro; IPR003598; Ig_c2.

InterPro; IPR003598; Ig_like.

InterPro; IPR003006; Ig_like.

InterPro; IPR003585; Neurexin-like.

Pfam; PF00047; 1g; 2.

SMART; SM00294; 4.1m; 1.

SMART; SM00410; IG_like; 1.
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Eukaryota; Metazoa; Chordata;
Eukaryota; Eutheria; Rodentia;
NCBI_TaxID=10090;
[1]
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01-MAY-1999 (TrEMBLrel.
01-JUN-2002 (TrEMBLrel.
Nectin-like protein 2.
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InterPro; IPR003598; Ig_c2.
InterPro; IPR003600; Ig_like.
InterPro; IPR003060; Ig_MHC.
InterPro; IPR003585; Neurexin-like.
Pfam; PP00047; ig; 2.
SMART; SM00294; 4.1m; 1.
SMART; SM00294; 1.1m; 1.
                                                                                                                                                                                                                                                Mus musculus (Mouse).
Eukaryota; Metazoa; C
Mammalia; Eutheria; R
NCBI_TaxID=10090;
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Q9QYL4;
Q1-MAY-2000
01-MAY-2000
01-JUN-2002
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IGSF4 OR
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Submitted (DEC-1998) to the
EMBL; AB021966; BAA87916.1;
                                                                                                                  MGD;
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Sciurognathi; Muridae;
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Adhesion protes...
IGSF4 OR RA175A.
Mus musculus (Mouse).
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InterPro; IPRO03598; Ig_c2.

InterPro; IPRO03600; Ig_11ke.

DR InterPro; IPRO03006; Ig_MHC.

DR Pfam; PF00047; 1g; 2.

DR SMART; SM00294; 4.1m; 1.

SMART; SM00294; 4.1m; 1.

YR SMART; SM00408; IGC2; 1.

YR Immunoglobulin domain.

SEQUENCE 295 **
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InterPro; IPR003585; Neurexin
Pfam; PF00047; Ig; 2.
SMART; SM00294; 4.1m; 1.
SMART; SM00408; IGc2; 1.
SMART; SM00410; IG_like; 1.
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01-MAY-2000 (TrEMBLrel. 1
01-JUN-2002 (TrEMBLrel. 2
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Eukaryota; Metazoa; Chordata;
Mammalla; Eutherla; Rodentia;
NCBI_TaxID-10090;
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Momoi T.;
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SMART; SM00294; 4.1m; 1.
SMART; SM00408; IGC2; 1.
SMART; SM00410; IG_like; 1.
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                  GCCATGGCCAGCAAGCCAGCGACGACCATCAGGTGGTTCAAAAGGGAACAAGGAACTCAAA
                                                                       ACCATTGGGGCAGTGGACCAC
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01-JUN-2002
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"Identification of a murine ortholog of the TS
Submitted (OCT-2001) to the EMBL/GenBank/DDBJ
EMBL; AY059394; AAL29692.1;
                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A. STRAIN=129/SVJ;
                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Mammalia; Eutheria; NCBI_TaxID=10090;
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  Vertebrata;
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  Euteleostomi;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pfam; PF00047; 1g; 3.
SMART; SM00294; 4.1m; 1.
SMART; SM00408; IGC2; 1.
SMART; SM00410; IG_like;
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submitted (AUG-1998) to the EMBL/GenBank/DDBJ
EMBL; AC005525; AAC32740.1; -.
HSSP; P80748; ZLOI.
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"Sequence analysis of a 2.5 mb region in 19q13.2 containing a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Immunoglobulin domain.
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NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GGACAGAATCTGTTTACTAAAGACGTGACAGTGATTGAAGGAGAAGTGGCAACCATCAGC
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                      CAGCTGATGCTGAAGGTGCACAAGGAGGACGACGGGGTCCCCGGTGATCTGCCAGGTGGAG
                                                                     LysGluLeuLysGlyValSerSerSerGlnGluAsnGlyLysValTrpSerValAlaSer
                                                                                            AAGGAACTCAAAGGCAAATCAGAGGTGGAGGAGTGGTCGGACATGTACACTGTGACCAGT
                                                                                                                                                                  LeuSerCysLeuValProArgSerArgProAlaAlaThrLeuArgTrpTyrArgAspArg
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; IPR003598; Ig_c2.

; IPR03600; Ig_like.

; IPR003006; Ig_MHC.

; IPR003585; Neurexin-like.
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                                                             Pfam; PF00047; 19; 3.
SMART; SM00409; IG; 3.
SMART; SM00408; IGc2; 3.
SMART; SM00410; IG_like; 2.
                                                                                                                                                                                                                                                                                                                                                           Fukami T., Maruyama T., Murakami Y.;
"Identification of a murine ortholog of the TSLC1-like of Submitted (OCT-2001) to the EMBL/GenBank/DDBJ databases EMBL; AF195562; AAG35584.1;
EMBL; AY059393; AAL29691.1;
MGD; MGI:2137858; Nec11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MECLI OR TSLL..
NECLI OR TSLL..
Mus musculus (Mouse).
Mus musculus (Motazoa; Chordata; Arvota; Metazoa; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
STRAIN=129/SVJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A. Zhou Y., Huang X., Yuan "Cloning and expression membrane protein.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Nectin-like protein 1 (Membrane glycoprotein).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (OCT-1999) to the
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; IPR003600; Ig_like.
; IPR003006; Ig_MHC.
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Sciurognathi; Muridae;
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                                                                                           ACAACTATCCCTCCTCCCACAACAACCACCACC
                                              AlaThrSerAsnMetGlySerTyrThrAlaTyrPheThrLeuAsnValAsnAspProSer
                                                                     ValTrpValLysGluGlySerGluProProLeuLysMetThrGlnGluSerAlaLeuIle
                                                                                                                                                                      ACTTGGGTGAGAGTCGATGATGAAATGCCT---
                                                                                                                                                                                           {\tt GluGlyGlnLysLeuLeuHisCysGluGlyArgGlyAsnProValProGlnGlnTyr}
                                                                                                                                                                                                                     GAAGGGGATGCATTTGAGTTAACGTGTGAAGCCATCGGGAAGCCCCAGCCTGTGATGGTA
                                                                                                                                                                                                                                            ValLeuTyrThrProThrAlaMetIleArg-----ProGluProAlaHisProArg
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                                                                                                                                                                                                                                                                                             {\tt CysSerValAsnHisGluSerLeuLysGlyAlaAspArgSerThrSerGlnArgIleGluscher} \\
                                                                                                                                                                                                                                                                                                                      TGCCAGGTGGAGCACCCTGCGGTCACTGGA---AACCTGCAGACCCAGCGCTATCTAGAA
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
BK134P22.1 (Novel protein similar to mouse IMMUNOSUPERFAMILY protein
BL2) (NECTIN-11ke protein 1).
BK134P22.1 OR NECL1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pfam; PF00047; 1g; 3.
SMART; SM00294; 4.1m; 1.
SMART; SM00408; IGc2; 1.
SMART; SM00410; IG_11ke; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Zhou Y., Du G., Yuan J., Qlang B.;
Submitted (APR-2001) to the EMBL/GenBank/DDBJ
EMBL; AL035403; CAB56227.1; -
EMBL; AF062733; AAD17540.2; -.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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InterPro; IPR003600; Ig_like.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003585; Neurexin-like.
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SerSerGlySerLysProAlaAlaArgLeuThrTrpArgLysGlyAspGlnGluLeuHis
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                           GCCATGGCCAGCCAGCCACCACCATCAGGTGGTTCAAAGGGAACAAGGAACTCAAA
                                                       IleIleThrGlyTyrLysSerSerLeuArgGluLysAspThrAlaThrLeuAsnCysGln
                                                                                 ATGATCGATATCCAGAAAGACACGGCAGTTGAAGGGGAGGAGGATTGAAGTCAACTGTACT
                                                                                                             ThrMetProValArgThrAlaLysSerLeuValThrValLeuGlyIleProGlnLysPro
                                                                                                                                       ACGGACCCCCCACAGGAGAGTTACACCACCATCACAGTCCTGGTTCCTCCACGTAACTTG
                                                                                                                                                                   SerIleSerIleSerAsnValAlaLeuAlaAspGluGlyGluTyrThrCysSerIlePhe
                                                                                                                                                                                                                           GluLysArgAlaLeuArgAspAsnArgIleGlnLeuValThrSerThrProHisGluLeu
                                                                                                                                                                                                                                                      GACTTCAGGCCTTTGAAGGACAGCAGGTTTCAGCTGCTGAATTTTTCTAGCAGTGAACTC
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AA; 47
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                              RA Kawai J. Shinagawa A. Shibata K. Yoshino M., Itoh M., Ishii Y.,
RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
RA Arakawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
RA Kuchl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuchl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuchl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Schrimi L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,
RA Schrimi L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Lee N.H.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
RA Lyons P., Marchionni L., Mashima J., Mazzarelli J., Mombaerts P.,
RA Lyons P., Marchionni L., Mashima J., Mazzarelli J., Mombaerts P.,
RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,
RA Suzuki H., Toyo-oka K., Wang K.H., Weitz C., Whittaker C., Wilming L.,
Wynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Q9D006;
01-JUN-2001 (TrEMBLrel. 1
01-JUN-2001 (TrEMBLrel. 1
01-DEC-2001 (TrEMBLrel. 1
2610301B19RLk protein.
                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
STRAIN-C57BL/6J; TISSUE-EMBRYO;
MEDLINE-21085860; PubMed-11217851;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
                      Wynshaw-Boris A.,
Hayashizaki Y.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus (Mouse).
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    Functional annotation of a
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EMBL, AKO11949; BABZ7933.1; -.
MGD; MGI:1914402; 2610301B19Rik.
InterPro; IPR003599; Ig.
InterPro; IPR003500; Ig_like.
InterPro; IPR003006; Ig_MHC.
Pfam; PF00047; Ig; 2.
SMART; SM00409; IG; 1.
SMART; SM00410; IG_11ke; 1.
SSQUENCE 549 AA; 60703 MW; 327
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16 GGACAGAATCTGTTTACTAAAGACGTGACAGTGATTGAAGGAGAAGTGGCAACCATCAGC
                                                                                                                                                                                                                                                                                                                                                                                      {\tt AspAlaAsnProProProPheLysSerValTrpSerArgLeuAspGlyGlnTrpProAsparation}
                                                                                                                                                                ThrArgPheAlaArgGlyArgArgIleThrCysValValLysHisProAlaLeuGluLys
                                                                                                                                                                                             CACAAGGAGGACGACGGGTCCCGGTGATCTGCCAGGTGGAGCACCCTGCGGTCACTGGA
                                                                                                                                                                                                                                       TCAGAGGTGGAGGAGTCGGACATGTACACTGTGACCAGTCAGCTGATGCTGAAGGTG
                                                                                                                                                                                                                                                                            ThrGlyLysProValAlaGlnIleAspTrp---GluGlyAspLeuGlyGluArgGluPhe
                                                                                                                                                                                                                                                                                                                                                            CAGAAAGACACGGCAGTTGAAGGG-----GAGGAGATTGAAGTCAACTGTACTGCCATG
                                                                                                          GCCAGCAAGCCAGCGACCATCAGGTGGTTCAAAGGGAACAAGGAACTCAAAGGCAAA
                                                                                                                                                                                                                                                                                                                                  GlyProAspSerSerIleAspGlyGlyAsnGluThrValAlaAlaValCysValSerSer
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SerThrGlnThrValAlaValHisHisProGlnTyrGlyPheSerValGlnGlyAspTyr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TGC-----CAGGTCAATAAGAGTGACGACTCAGTGATC-----CAGCTCCTGAACCCC
                           ATCGGGAAGCCCCAGCCTGTGATGGTAACTTGGGTGAGAGTCGATGATGAAAATGCCTCAA 753
                                                      TyrAspGlyAsnTrpPheValGlyArgLysGly
                                                                                                                                      AACCTGCAGACCCAGCGCTATCTAGAAGTGCAGTATAAACCGCAAGTGCATATCCAGATG
                                                                                                                                                                                                                       SerThrIleSerPheLeuAsnGluThrAlaThrIleValSerGlnTyrGluLeuPhePro
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                                                                                 ACTTACCCTCTGCAAGGCCTAACCCGGGAAGGGGATGCATTTGAGTTAACGTGTGAAGCC
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ProAlaAspValGlnAspIleAlaThrGluHisLysLysLeuProPheProLeuSerThr
                                                                                       TyrIleSerAspProProThrThrThrThrLeuGlnProThrValGlnTrpHisSerSer
                                                                                                    GlyValTyrValCysLysValSerAsnSerLeuGlyGlnArgSerAspGlnLysValIle
                                                                                                                                                          GGTACTTACCGCTGTGAGGCTTCCAACATAGTGGGAAAGGCTCATTCGGACTATATGCTG
                                                                                                                                                                                {\tt GlyLeuLeuAlaSerAspAsnThrLeuHisPheValHisProLeuThrValAsnTyrSer}
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Search completed: November 20, 2002, 07:56:07 Job time: 77.25 secs

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     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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2: /cgn2_6/ptodata/1

3: /cgn2_6/ptodata/1

4: /cgn2_6/ptodata/1

5: /cgn2_6/ptodata/1

6: /cgn2_6/ptodata/1
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/cgn2_6/ptodata/1/18a/BCTUS_COMB.pep:*
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US-08-429-742-2
US-08-429-742-3
US-08-6217-299-3
US-08-602-725-3
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172	172	172	172.5	172.5	172.5	173.5	174.5	174.5	174.5	177	177	179.5	179.5	183.5	183.5	183.5	185
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US-09-991-326-12	US-09-707-802-12	US-08-752-307B-12	US-09-077-955-33	US-08-644-271-29	US-08-374-834-16	US-09-724-864-62	PCT-US95-08493-21	PCT-US95-08493-19	PCT-US95-08493-15	PCT-US94-05277-2	US-09-041-886-25	US-09-135-080-8	US-08-414-657D-60	US-09-540-245A-17	US-08-684-594-3	US-08-432-016-3	US-08-414-657D-45
Sequence 12, Appl	12,	Sequence 12, Appl	ω 33',	Sequence 29, Appl	-	Sequence 62, Appl	-	-	•		Sequence 25, Appl		Sequence 60, Appl	Sequence 17, Appl	Sequence 3, Appli	Sequence 3, Appli	Sequence 45, Appl

#### ALIGNMENTS

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; TOPOLOGY: 1; MOLECULE TYPE: US-08-659-984A-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: Two Embarcadero Ctr., 8th Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
COURTERNT APPLICATION DATA:
Query Match
Best Local Similarity
Matches 137; Conserv
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                                                                                                                                                                                             TELEFAX: 415-326-2422
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 15
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Heslin, James M.
REGISTRATION NUMBER: 29,
                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/485,152
FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Jacobson-Croak, Kirsten L.
TITLE OF INVENTION: Assays for Detecting Beta-Secretase
TITLE OF INVENTION: Inhibition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Anderson, John APPLICANT: Sinha, Sukanto APPLICANT: Jacobson-Croak,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/659,984A FILING DATE: 07-JUN-1996 CLASSIFICATION: 436
                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                 LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
                                                                                                                                                          amino acid
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Two Embarcadero Ctr., 8th Floor
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    Conservative
                                                                                                                        linear
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                      35.4%; Score 625.5; DB 1
39.0%; Pred. No. 1.2e-47;
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    66;
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    Mismatches
                                         DB 2;
    121;
                                         Length 421;
    Indels
    27;
    Gaps
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Query Match
Best Local Similarity
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                                                                                                                                                                    TELEFAX: 415-326-2422 INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
APPLICANT:
                                                                                                                                                                                               NAME: Heslin, James M.
REGISTAN NUMBER: 29,541
REFERENCE/DOCKET NUMBER: 15
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/480,498
FILING DATE: 07-JUN-1995
ATTORREY/AGENT IMFORMATION:
NAME: 1200115 Targot
                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                          MOLECULE TYPE:
                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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CITY: S
                                                                                       STRANDEDNESS: SIL
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Two Embarcadero Ctr., 8th Floor
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Sinha, Sukanto
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              35.4%;
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ER: 15270-002210US
              Score
Pred.
               No.
                              625.5;
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               1.2e-47;
                              DВ
                              4.
                              Length 421;
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US-08-659-984A-5
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                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/659,984A
FILING DATE: 07-JUN-1996
CLASSIFICATION: 436
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/485,152
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
APPLICANT:
TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                              NAME: Healin, James M.
REGISTRATION NUMBER: 29,541
REFERENCE/DOCKET NUMBER: 155
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Townsend
                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      307
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     300 PPPTTTTTTTTTTTILTIIT-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             242 WVRVDDEM--PQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTI 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 190 RVDHESLNATPQVAMQVLEIHYTPSVKI---IPSTPFPQEGQPLILTCESKGKPLPEPVL 246
                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM:
SOFTWARE: PatentI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET:
                                    STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       66
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                                                                          LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVEGEEIE 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LPTTIIPSLTTATVTTTVAITTSPTTSATTSSIRDPNALAGQNGP----DH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WTKDGGELPDPDRMVVSGRELNILFLNKTDNGTYRCEATNTIGQSSAEYVLIVHDVPNTL 306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    QVEHPAVTGNLQ-TQRYLEVQYKPQVHIQMTYPLQGLTREGDAFELTCEAIGKPQPVMVT 241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            VNCTAMASKPATTIRWFKGNKELKGKSEVEEWS - - - DMYTVTSQLMLKVHKEDDGVPVIC 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GQNLFTKDVTVIEGEVATISCQVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQLLNFS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    San Francisco
California
                                                     amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Townsend and Townsend and Crew Two Embarcadero Ctr., 8th Floor
                                                                      444 amino acids
               linear
                                                                                                                                                                                                                                                                                                                                                                                      IBM PC compatible SYSTEM: PC-DOS/MS-DOS PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Jacobson-Croak, Kirsten L. VENTION: Assays for Detecting
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                                                                                                                                                  415-326-2400
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OM protein - protein search, using sw model
                                                                                       GenCore version 5.1.3 Copyright (c) 1993 - 2002 Compugen Ltd.
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Run on: November 20, 2002, 07:34:43; Search time 15 Seconds (without alignments) 2153.411 Million cell updates/sec

Title: Perfect score: Sequence:

US-09-778-187B-4\_COPY\_21\_356
1765
1 IPTGDGQNLFTKDVTVIEGE.....LTIITDSRAGEEGTIGAVDH 336

Scoring table: BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched: 283224 seqs, 96134422 residues

Total number of hits satisfying chosen parameters: 283224

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR\_73:\*
1: pir1:\*
2: pir2:\*
3: pir3:\*
4: pir4:\*

Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	8	7	o,	ហ	4	ω	2	1	NO.	Result
205.5	205.5	206	206.5	206.5	207.5	209.5	209.5	210	212	214	215	218	218	220	220	220	220	222	222	222	223	226	228.5	234	238	245	261.5	261.5	Score	
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932	812	847	868	862	345	584	345	518	4391	1011	478	417	392	538	467	417	392	1092	725	530	344	4162	7962	1088	725	407	5198	5175	Length	
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cell adhesion mole	cell adhesion mole		CD22 homolog/B lym		opioid-binding cel	s-gicerin precurso	opioid-binding pro	poliovirus recepto	perlecan precursor	lin	PRR2 alpha - human		poliovirus recepto	PRR2 delta - human	poliovirus recepto	poliovirus recepto	н	cell	neural cell adhesi	poliovirus recepto	neurotrimin - rat	connectin/titin -	c titi	cell	8	hypothetical prote	н	al pro	Description	

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231 AIGKPQPVMVTWVRVDDEMPQHAVL----SGPNLFINNLNKTDNGTYRCEASNIVGKAHS 286

## ALIGNMENTS

Db 2355 EGDAGKYTCEADNSAGSVEQDVNVNVITIPKIEKDGIPSDYESQQNERVVISCP 2408	
QY 174 EDDGVPVICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDAFELTCE 230	
Db 2310 CPA-TGKPEPDITWFKDGEAIHIENIADIIPNGELNGNQLKITRIK 2354	
Qy 128 CTAMASKPATTIRWFKGNKELKGKSEVEEWSDMYTVTSQLMLKVHK 173	
Db 2250 RLTILNVTENDEGQYSCRVKNDAGENSFDFKATVLVPPTIIMLDKDKNKTAVEHSTVTLS 2309	
Qy 70 KVSLTNVSISDEGRYFCQLYTDPPQESYT-TITVLVPPRNLMIDIQKD-TAVEGEEIEVN 127	
Db 2200 VTAIKGGALPEKCPIDDDKNFKGQIIWLRNYQPIDLEAEDARITRLSNDR 2249	
Qy 14 VTVIEGEVATISCOVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQLLNFSSSEL 69	
Query Match 14.8%; Score 261.5; DB 2; Length 5175; Best Local Similarity 24.6%; Pred. No. 8.2e-10; Matches 87; Conservative 65; Mismatches 130; Indels 71; Gaps 15;	
A;Map position: X A;Map position: X A;Map position: X A;Thorons: 85/1; 120/1; 334/3; 370/1; 477/2; 606/3; 664/1; 935/3; 977/1; 1051/3; 1184 A;Introns: 85/1; 120/1; 334/3; 370/1; 477/2; 606/3; 664/1; 935/3; 977/1; 1051/3; 1184 A;Introns: 85/1; 2699/3; 2759/1; 2852/1; 2889/3; 2913/3; 2941/1; 2967/3; 2991/3; 303 i; 2512/2; 2593/3; 2699/3; 2759/1; 2852/1; 2889/3; 2913/3; 2941/1; 2967/3; 2991/3; 303 i; 4225/1; 4361/1; 4408/1; 4456/1; 4498/1; 4647/3; 4838/1; 4879/1; 4941/1; 5011/1; 50	
C;Genetics: A:Gene: CESP:F15G9.4a	
A;Cross·references: EMBL:247070; PIDN:CAA87344.1; GSPDB:GN00028; CESP:F15G9.4a A;Experimental source: clone T09B9	
A; Molecule type: DNA A; Residues: 1-5175 <wi2></wi2>	
A; Status: preliminary; translated from GB/EMBL/DDBJ	
ce number: 219929 on: T24733	
R;Kershaw, J. submitted to the EMBL Data Library, December 1994	
A; Experimental source: clone F15G9	
A;Status: preliminary; translated from GB/EMBL/DDBJ	
R;Sulston, J. submitted to the EMBL Data Library, December 1994	
C;Date: 15-Oct-1999 #text_change 18-Feb-2000 C;Date: 15-Oct-1999 #text_change 18-Feb-2000 C;Accession: T20992; T24733	
hypothetical protein F15G9.4a - Caenorhabditis elegans	
RESULT 1 T20992	

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Molecule type: DNA
A; Residues: 1-5198 <WI2>
A; Cross-references: EMBL: Z47070;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               submitted to the EMBL Data Library, A; Reference number: Z19929
A; Accession: T24734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Molecule type: DNA
A;Residues: 1-5198 <WIL>
A;Cross-references: EMBL:Z47068; PIDN:CAA87336.1; GSPDB:GN00028; CESP:F15G9.
A;Experimental source: clone F15G9
R;Kershaw, J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hemicentin precursor - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 11-Jan-2000 #sequence_revision 11-Jan-2000 C;Accession: T43290; T20993; T24734
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A; Residues: 1-5198 < VOG>
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Best Local
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   DIMLYVYDPPTTIPP---
                                       VYARP-PAKITWLKAGKPLQSDKFVKTSANGQKLYLFKLRETDSSKYTCIATNEAGTDKR
                                                                   AIGKPQPVMVTWVRVDDEMPQHAVL----SGPNLFINNLNKTDNGTYRCEASNIVGKAHS
                                                                                                                                                                                                                        CTAMASKPATTIRWFKG---
                                                                                                                                                                                                                                                                                          KVSLTNVSISDEGRYFCQLYTDPPQESYT-TITVLVPPRNLMIDIQKD-TAVEGEEIEVN 127
                                                                                                                                                                                                                                                                                                                                VTAIKGGALPFKCPID--DDK-----NFKGQIIWLRNYQPIDLEAEDARITRL---SNDR
                                                                                                                                                                                                                                                                                                                                                                   VTVIEGEVATISCQVNKSDDSVIQLLNPNRQTIYFRDFRPL----KDSRFQLLNFSSSEL
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                                                                                                           EGDAGKYTCEADNSA--GSVEQDVNVNVITIPKIEKDGIPSDYESQ----QNERVVISCP
                                                                                                                                             EDDGVPVICQVEHPAVTGNLQTQRYLEVQYKPQVH----IQMTYPLQGLTREGDAFELTCE 230
                                                                                                                                                                                  CPA-TGKPEPDITWFKDGEAIHIENIADIIPNGELNG-------NQLKITRIK
                                                                                                                                                                                                                                                        RLTILNVTENDEGQYSCRVKNDAGENSFDFKATVLVPPTIIMLDKDKNKTAVEHSTVTLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DFKVSMLVAPSFDEPNIVRRITVNSGNPSTLHCPAKGSPSPTITWLKDGNAIE 2520
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Hemicentin is require
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                                                                                                                                                                                                                                                                                                                                                                                                                      14.88;
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Pred. No. 8.3e-10;
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   PTTTTTTTTTTTLTITTDSRAGE
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nypotnetical protein DKFZp566B0846.1 - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change
C;Accession: T08732
R;Ottenwaelder, B.; Obermaier B. Va....
                                                                                                                                             C;Comment: This protein mediates and regulates various cell-cell interactions C;Superfamily: neural cell adhesion molecule; fibronectin type III repeat homo F;413-475_Domain: immunoglobulin homology <TMM>F;512-589_Domain: fibronectin type III repeat homology <3FR>
                                                                                                                                                                                                                                                                                                                                                        R;Kudo, M.; Takayama, E.; Tadakuma, T.; Shiokawa, K. Biochem. Biophys. Res. Commun. 245, 127-132, 1998 A;Title: Molecular cloning of ssd-form neural cell adhesion A;Reference number: JE0099; MUID:98204770; PMID:9535795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              neural cell adhesion molecule 1 - African clawed frog
N;Alternate names: N-CAM 1
C;Species: Xenopus laevis (African clawed frog)
C;Date: 19-May-1998 #sequence_revision 29-May-1998 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: EMBL:AL050071
A;Experimental source: fetal kidney;
C;Genetics:
A;Note: DKFZp566B0846.1
                                                                                                                                                                                                                                                                 A;Cross-references: DDBJ:AB008162; NID:g3116226; PIDN:BAA25931.1; PID:g3116227
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May 1999
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              Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.
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Gapop 10.0 , Gapext 0.5
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           SwissProt_40:*
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        GenCore version 5.1.3
(c) 1993 - 2002 Compugen Ltd
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CD22_HUMAN
CD22_HUMAN
OPCM_BOVI
NPHN_MOUSE
CEPU_CHICK
OPCM_HUMAN
NEO1_HUMAN
NEO1_HUMAN
NEO1_CHICK
OPCM_RAT
LAMP_CHICK
OPCM_RA
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PVR2_MOUSE
NCA2_XENLA
G55A_CHICK
PVR_CERAE
PVR_CERAE
PVR2_HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SUMMARIES
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T mus musculu
5 xenopus lae
2 gallus gall
6 cercopithec
2 homo sapien
1 homo sapien
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mus musculu
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8888		RESULT NCA1_X ID NA AC P DT 0										
Amp Amp	01-AP 15-JU Neura 180).	RESULT 1 NCA1_XENLA ID NCA1_ AC P1617 DT 01-AP		44	4 4 ω 2	410	39	37 8	36	<b>ω</b>	34	
Xenopus laevis (Afri Eukaryota; Metazoa; Amphibia; Batrachia;	01-APR-1990 15-JUN-2002 Neural cell 180).	T 1 XENLA NCA1_XENLA P16170; 01-APR-1990		185.5 182.5	185.5 185.5	187.5 185.5	188	189	189	189.5	191.5	
vis (Afi Metazoa atrachia	(Rel. 14 (Rel. 41 adhesion	STAI		10.5 10.3	10.5 10.5	10.6	10.7	10.7	10.7	10.7	10.8	
rican ; Chor a; Anu	a``	STANDARD;		1241 2012	764 1234	ა ა ა ა 8	1447	588 858	853	646	583	
cla dat ra;	st ecu	D; Created)		11			ь,			_	1	
Norma: Xenopus laevis (African clawed frog). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleost Eukaryota; Metazoa; Chordata; Mesobatrachia; Pipoidea; Pipidae; Manphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;	, Last sequence update) , Last annotation update) molecule 1, 180 kDa isoform p	PRT; 1088 AA.	ALIGNMENTS	NPHN_HUMAN DSCA_HUMAN	ICCR_DROME NPHN_RAT	LAMP_RAT	DCC_MOUSE	NCA1_RAT	NCA1_BOVIN	MU18_HUMAN	C166_HUMAN	
; Euteleostomi; a; Pipidae;	precursor (N-CAM			_	Q08180 drosophila Q9r044 rattus norv	Q13449 homo sapien Q62813 rattus norv		P13596 rattus norv	-	homo	Q13740 homo sapien	

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SM00408; IGc2;	SMART; SM00060; FN3; 2.		PE00041;	Pro; IPROUSSY8	TPRO03000;	TPROODER,	•		PIR: S09600: IJXLNL.	EMBL; M25696; AAA49909.1;		license	http://	ified and this statement is not removed. Usage by and for commer	use by non-profit institutions as long as its content is in no way	the European Bioinformatics Institute. There are no restrictions on its	between the Swiss Institute of Bioinformatics and the EMBL outstation -	This SWISS-PROT entry is copyright. It is produced through a collaboration		CONTAINS 2	-!- SIMILARITY: CONTAINS 5 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS.	-!- SIMILARITY: BELONGS TO THE IMMUNOGLOBULIN SUPERFAMILY.	EARLY NEURAL DEVELOPMENT.				-!- TISSUE SPECIFICITY: EXPRESSED IN NEURON AND IN PRESUMPTIVE NEURAL	g.	-!- ALTERNATIVE PRODUCTS: 2 isoforms; N-CAM 180 (shown here) and	-!- SUBCELLULAR LOCATION: Type I membrane protein.	NEURITES, ETC.	NEURON-NEURON ADHESION, NEURITE FASCICULATION, OUT		Nucleic Acids Res. 17:10321-10335(1989).	molecule (NCAM).";		lopmenta	Krieg P.A., Sakaguchi D.S., Kintner C.R.;	MEDLINE-90098871; PubMed-2481269;	SEQUENCE FROM N.A.	[1]	NCBI_TaxID=8355;	Xenopodinae; Xenopus.	Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;

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RESULT 2
PVRI_PIG
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Best Local
PVRI_PIG STANDARD; PRT; 51: Q9GL76; 16-CCT-2001 (Rel. 40, Created) 16-CCT-2001 (Rel. 40, Last sequence upda 15-JUN-2002 (Rel. 41, Last annotation up Poliovirus receptor related protein 1 primediator C) (HveC) (Nectin 1). PVRL1 OR PRR1 OR HVEC. Sus scrofa (Pig). Eukaryota; Metazoa; Chordata; Craniata;
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4 1049
8 AA; 117778
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PROBAB
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IG-LIKE C2-TYPE DOMAIN 2.
IG-LIKE C2-TYPE DOMAIN 2.
IG-LIKE C2-TYPE DOMAIN 3.
IG-LIKE C2-TYPE DOMAIN 3.
IG-LIKE C2-TYPE DOMAIN 4.
IG-LIKE C2-TYPE DOMAIN 5.
FIBRONECTIN TYPE-III 2.
FIBRONECTIN TYPE-III 2.
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2; Mismatches
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    Craniata;
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    Vertebrata;
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.9e-10;
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   TNVSISDEGRYFCQLYTDPP-
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SMART; SM00409; IG; 3.
SMART; SM00410; IG_like;
SMART; SM00408; IGc2; 2.
SMART; SM00406; IGv; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
MEDLINE=21176378; PubMed=11277703;
Milne R.S.B., Connolly S.A., Krumm
Cohen G.H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                            modified and this statement is not rementities requires a license agreement
                                                                                                                                                                                                                                                                                                                                                                                                                                     or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mammalia; Eutheria; NCBI_TaxID=9823;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 a functional alphaherpesvirus rology 281:315-328(2001).
- FUNCTION: PROBABLY INVOLVED 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SUBUNIT: Interacts with HSV glycoprotein D (gD) (By similar SUBCELLULAR LOCATION: Type I membrane protein. SUPERFAMILY. SIMILARITY: BELONGS TO THE IMMUNOGLOBULIN SUPERFAMILY. SIMILARITY: CONTAINS 1 IMMUNOGLOBULIN-LIKE V-TYPE DOMAIN. SIMILARITY: CONTAINS 2 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ALPHAHERPESVIRUS (HSV-1,
                VTVIEGEVATISCOVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQLLNFSSSELKVSL
                                                                                                                                                                                                                                                                                                                                                                                                                  AF308632; AAG30281.1;
                                                                                                                                                                                                                                                                                       Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                non-profit institutions as long
                                                                                                                                                                                                                                                                                                                                                                                      IPR003599;
IPR003006;
                                                                                                                                                                                                                                                                                                                                                         IPR003596;
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IPR003600;
                                     Conservative
                                                                                                                                                                                                                                                                                                 Immunoglobulin
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Ig_like.
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                                             . 4 %;
                                                                           MW;
                                     49;
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VAIYNPAMGVSVLAPYR---
                                                                                                                                                                                                                              POTENTIAL.

POLICYRUS RECEPTOR RELAT
EXTRACELLULAR (POTENTIAL)
POTENTIAL.

CYTOPLASMIC (POTENTIAL).

IG-LIKE V-TYPE DOMAIN.
                                            Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      is not removed.
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N-LINKED
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O1-JUN-2002 (TrEMBLrel. 21, Created)
O1-JUN-2002 (TrEMBLrel. 21, Last sequence update)
O1-JUN-2002 (TrEMBLrel. 21, Last annotation update)
O1-JUN-2002 (TrEMBLrel. 21, Last annotation update)
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"Identification of murine orthologue of the TSLC1 gene.";
Submitted (OCT-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AF434663; AAL86736.1;
SEQUENCE 445 AA; 48664 MW; C5D5A070DAF70E55 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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Sciurognathi;
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thi; Muridae; Murinae; Mus
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099412 homo sapien
09v490 drosophila
073633 xenopus lae
095727 homo sapien
010465 homo sapien
010465 homo sapien
08wz42 homo sapien
08wz42 homo sapien
098918 gallus gall
073634 xenopus lae
099121 homo sapien
09046f gallus gall
091vf9 mus musculu
098x4 brachydanio
099x94 brachydanio
09up10 mus musculu
09ue16 homo sapien
096bj1 homo sapien
096bj1 homo sapien
096bj4 homo sapien
096bj4 homo sapien
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01-JUN-2002
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RA175.
                                 Q9CRY3 PRELIMINARY; PRT; 494 AA.
Q9CRY3;
01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation updata)
3100001108Rik protein (Fragment).
IGSF4 OR 3100001108RIK.
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                     Mus musculus (Mouse)
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EMBL; AB064265; BAB8
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RA Alzawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Alzawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Alzawa K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
RA Kiehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kiehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kiehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kiehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kiehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kiehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kaeli K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
Ra Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
Lyons P., Marchionni L., Mashima J., Mazzarelli J., Mombaerts P.,
RA Mordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,
RA Aurschi-M., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,
RA Sasaki H., Sato K., Wang K.H., Weitz C., Whittaker C., Wilming L.,
Wynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
RA Hayashtzaki Y.,
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InterPro; IPR003006; Ig_MH
Pfam; PF00047; Ig; 3.
SWART; SW00409; IG; 3.
SWART; SW00408; IGc2; 3.
SWART; SW00410; IG_like; 1
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InterPro;
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EMBL; AK013911; BAB29050.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      494
                  PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ig_c2.
Ig_like.
Ig_MHC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99.1%;
96.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 1749.5; DB 11; Pred. No. 1.7e-137;
                  PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      F5E09ABA1857ABC0 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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Best [
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"Cloning of a novel human cDNA encoding a superfamily.", AF69029.1; -.
Submitted (MAR-1999) to the EMBL/GenBank/D EMBL; RP132811; AAF69029.1; -.
InterPro; IPR003599; Ig.-C2.
InterPro; IPR003599; Ig.-C2.
InterPro; IPR003500; Ig_like.
InterPro; IPR003500; Ig_like.
InterPro; IPR003006; Ig_MHC.
Pfam; PF00047; 1g; 3.
SMART; SM00409; IG; 3.
SMART; SM00409; IG; 3.
SMART; SM004009; IGC2; 3.
Inmunoglobulin domain.
SEQUENCE 442 AA; 48537 MW; 68183E32387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                           Q9D6E7 PRELIMINARY;
Q9D6E7;
Q1-JUN-2001 (TrEMBLrel. 1
Q1-JUN-2001 (TrEMBLrel. 1
Q1-DEC-2001 (TrEMBLrel. 1
2900073G06R1k protein.
IGSF4 OR 2900073G06R1K.
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01-JUN-2001
01-DEC-2001
          SEQUENCE FROM N.A
STRAIN-C57BL/6J; '
MEDLINE-21085660;
                                                             Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
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Mammalia; Eutheria;
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NECL2.
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                                                                                                                                                                                                                 PPTTTTTTTTTTTLTIITDSRAGEEGSIRAVDH
                                                                                                                                                                                                                                                                                                                                                                                       LLNFSSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQRDTAVE
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31; Conservative
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Metazoa; Chordata; C
~haria; Primates; (
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(TrEMBLrel.
(TrEMBLrel.
protein 2.
N.A.
6J; TISSUE-HIPPOCAMPUS;
5660; PubMed-11217851;
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98.5%;
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Last
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Pred. No. 1.3e-136;
2; Mismatches 3;
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                                                             Craniata; Ver
Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Craniata; Vertebrata; |
Catarrhini; Hominidae;
                                                                                                                                                              PRT;
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                                                                        Vertebrata;
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                                                               Muridae;
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                                                               Euteleostomi; 
; Murinae; Mus
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RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S., RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I., Ra Salto T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Salto R., RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T., RA Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H., RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J., RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J., RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T., RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G., RA Blake J., Beffelli D., Bojunga N., Carninci P., de Bonaldo M.F., RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M., RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H., RA Lyons P., Marchionni L., Mashima J., Wazzarelli J., Wambaerts P., Ra Sasaki H., Toyo-oka K., Wang K.H., Weitz C., Whittaker C., Wilming L., RA Wynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S., Turnal Sanctional 
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Q9Z2H8
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Q9Z2H8;
Q1-MAY-1999 (TrEMBLrel. 10
Q1-MAY-1999 (TrEMBLrel. 10
Q1-JUN-2002 (TrEMBLrel. 2
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SMART; SM00409; IG; 3.
SMART; SM00408; IGc2; 3.
SMART; SM00410; IG_like; 1.
Immunoglobulin domain.
SEQUENCE 336 AA; 37157 MW;
                                                   Mus musculus (Mouse)
Eukaryota; Metazoa; (
Mammalia; Eutheria; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR003599; Ig.
InterPro; IPR003598; Ig_c2.
InterPro; IPR003600; Ig_11ke.
InterPro; IPR003006; Ig_MHC.
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REMBL: AK013775; BAB28988.1; -.
                                                                                                                                                                                                                                    Q9Z2H8
                              NCBI_TaxID=10090
                                                                                                               IGSF4 OR NECL2
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nilarity 100.0%;
Conservative 0
                                                                                                                                                                                                                                    PRELIMINARY;
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                                                     Chordata;
Rodentia;
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Pred. No. 2.3e-120;
0; Mismatches 0;
                                                   Craniata; Vertebrata;
Sciurognathi; Muridae;
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                                                                       Euteleostomi;
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Matches 224
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Best Local :
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InterPro; IPR003600; Ig_like.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003585; Neurexin-like.
Pfam; PF00047; ig; 2.
SMART; SM00294; 4.1m; 1.
SMART; SM00408; IGc2; 1.
SMART; SM00410; IGc2; 1.
                                                                                                                      InterPro; IPR003006; Ig_MHC.
InterPro; IPR003585; Neurexin
Pfam; PF00047; ig; 2.
SMART; SM00294; 4.1m; 1.
SMART; SM00408; IGc2; 1.
SMART; SM00410; IG_like; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Adlic.
IGSF4 OR RAL/...
Mus musculus (Mouse).
Mus musculus (Mouse).
Metazoa; Chordata;
Metazoa; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q9QYL4;
                                                                                                                                                                                                                                                                                          Submitted (DEC-1998) to the EMEMBL; AB021966; BAA87916.1; -. MGD; MGI:1889272; Igsf4.
                                                                                                                                                                                                                                                                                                                                                                Momoi T.; "RA175, a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAY-2000 (TrEMBLrel. 13, 01-MAY-2000 (TrEMBLrel. 13, 01-JUN-2002 (TrEMBLrel. 21, Adhesion protein RA175C. IGSF4 OR RA175C.
                                                                                     Immunoglobulin SEQUENCE 306
                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A. Soyama A., Fujita
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (APR-1998) to the EMBL; AF061260; AAC67243.1; MGD; MGI:1889272; Igsf4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=10090;
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"Cloning of
                                                                                                                                                                                                                                                           InterPro;
                                                                                                                                                                                                                                                                             InterPro;
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  224;
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227; Conserv
                        Similarity
                                                                                                                                                                                                         IPR003598; Ig_c2.
IPR003600; Ig_like.
IPR003006; Ig_MHC.
IPR003585; Neurexin-like.
                                                                                                                                                                                                                                                                                                                                          novel neuron specific adhesion protein.", (DEC-1998) to the EMBL/GenBank/DDBJ datak
    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           68.2%; ilarity 100.0%; Conservative
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a novel cDNA
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AA; 32
                                                                                   domain.
AA; 3:
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                                                                                     33522 MW;
                        66.4%;
94.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yuan J., Qiang B.;
encoding a member of
the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                   Χ.,
    <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Last sequence update)
Last annotation update)
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                   Score 1171.5; DB Pred. No. 1.3e-89;
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Pred. No. 2.4e-92;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRT;
                                                                                     A4CE37B0F23554D5 CRC64;
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    Mismatches
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                                           DB 11;
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    11;
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Best Local S
Matches 217
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InterPro; IPR003600; Ig_like.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003585; Neurexin-like.
Pfam; PF00047; ig; 2.
SMART; SM00294; 4.1m; 1.
SMART; SM00408; IGc2; 1.
SMART; SM00410; IG_like; 1.
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01-MAY-2000 (TrEMBLrel. 1
01-JUN-2002 (TrEMBLrel. 2
Adhesion protein RA175A.
IGSF4 OR RA175A.
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IGSF4 OR RAL/...
Mus musculus (Mouse).
Mus-musculus (Mouse).
Thervota; Metazoa; Chordata;
Theria; Rodentia;
                                                                                                                                                                                                                                                                                                                            "RA175, a novel neuron specific Submitted (DEC-1998) to the EMBL EMBL; AB021964; BAA87914.1; -. MGD; MGI:1889277; 19sf4.
                                                                                                                                                                                                                                    SEQUENCE
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                                             EAIGKPQPVMYTWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYM
                                                                                                                     KVHKEDDGVPVICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDAFELTC
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                                                                                                                                                                                               217;
                                                                                                                                                                                               Similarity 95.0
17; Conservative
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                                                                                                                                                                                                                                              domain
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                                                                                                                                                                                                       Score 1151;
Pred. No. 6.
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Sciurognathi; Muridae;
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                                                                                                                                                                                                Mismatches
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Q9QYL5

PRELIMINARY;

PRT;

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RESULT
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Best Local S
Matches 205
         Soyama A., Fujita E., Urase K., Mu Momo! T.;
"RA175, a novel neuron specific ac submitted (DEC-1998) to the EMBL/C EMBL; AB021967; BAA87917.1; -. MCD; MCI:1889272; Igst4.
Interpro; IPR003598; Ig-c2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro: IPR003006; 19_MHC.
InterPro: IPR003585; Neurexin
Pfam: PF00047; 1g; 2.
SWART: SW00294; 4.1m; 1.
SWART: SW00408; IGC2; 1.
SWART: SW00410; IG_11ke; 1.
                                                                                                                                                  01-MAY-2000 (TrEMBLrel. 1
01-MAY-2000 (TrEMBLrel. 1
01-JUN-2002 (TrEMBLrel. 2
Adhesion protein RA175N.
IGSF4 OR RA175N.
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01-MAY-2000 (TrEMBLrel. 1
01-JUN-2002 (TrEMBLrel. 2
Adhesion protein Ral75B.
IGSF4 OR RA175B.
                                                                                                                                                                                              Q9QYL3;
Q9QYL3;
Q1-MAY-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Immunoglobulin domain. SEQUENCE 289 AA; 31811 MW;
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Momoi T.;
"RA175, a novel no
                                                                                                                   Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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                                                                                      SEQUENCE FROM N.A.
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InterPro; IPR003600; Ig_like.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003588; Neurexin-like.
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Ig_c2.
Ig_like.
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90.3%;
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EMBL/GenBank/DDBJ databases
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Last sequence update)
Last annotation updat
                                       Lric adhesion protein.",
EMBL/GenBank/DDBJ datah
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 1062; D
Pred. No. 1.6e
2; Mismatches
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                                                                                                                                                                                                                      PRT;
                                                                                                                     Craniata; Ver
Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8D1B836D0565AEA4
                                                                          Mukasa
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                                                                                                                                                                                                                      278
                                                                                                                              Vertebrata; Euteleostomi;
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                                                                         Kouroku
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                                                                                                                     Muridae;
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                                           databases
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                                                                                                                     Murinae;
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Matches 123; Conserv
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Best Local
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InterPro; IPR003585; Neurexin-11ke.
Pfam; PP00047; ig; 2.
SMART; SM00294; 4.lm; 1.
SMART; SM00408; IGC2; 1.
SMART; SM00410; IG_11ke; 1.
Immunoglobulin domain.
SEQUENCE 278 AA; 30636 MW; A295F
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01-JUN-2002 (TrEMBLrel.
01-JUN-2002 (TrEMBLrel.
01-JUN-2002 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fukami T., Maruyama T., Murakami Y.;
"Identification of a murine ortholog of the TSLC1-like gene
Submitted (OCT-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AY059394; AAL19692.1; -.
SEQUENCE 388 AA; 42723 MW; 8E3A9DF1C3B9D23E CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID-10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q8R464
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STRAIN=129/SVJ;
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                                                                                                                                                                                                                                                                                                       KVHKEDDGVPVICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDAFELTC
                                                                                                                                                                                                                                         FSSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVEGEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LYVYD------TTATTEPAVHDSRAGEEGTIGAVDH
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WVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIPP
                                      AQNQALPSGHSKQTQYVLDVQYSPTARIHAS---QAVVREGDTLVLTCAVTGNPRPNQIR
                                                                         VEHPAVTG--NLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDAFELTCEAIGKPQPVMVT
                                                                                                                                VELSCLVPRSRPAAVLRWYRDRKELKGVSSGQENGKVWSVASTVRFRVDRKDDGGIVICE
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88.1%;
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Last annotation updat
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Pred. No. 7.3e-44;
2; Mismatches 112;
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2. Lamerdin J.E., McCready P.M., Skowronski E., Adamson A.W.,

2. Lamerdin J.E., McCready P.M., Skowronski E., Adamson A.W.,

2. Lamerdin J.E., McCready P.M., Skowronski E., Adamson A.W.,

2. McCready P.M., Gardon L., Kyle A., Ramirez M., Stilwager

2. Lamerdin J., Poundstone P., Christensen M., Georgescu A., Avila

2. Lau S., Attix C., Andreise T., Trankheim M., Anico-Keller G.,

2. Lau S., Attix C., Andreise T., Trankheim M., Anico-Keller G.,

2. Coefield J., Duarte S., Lucas S., Bruce R., Thomas P., Quan G.,

2. McCoefield J., Duarte S., Lucas S., Bruce R., Thomas P., Quan G.,

2. Ak Kronmiller B., Arellano A., Montgomery M., Ow D., Nolan M., Tron

2. Ak Kobayashi A., Olsen A.S., Carrano A.V.,

2. Kobayashi A., Olsen A.S., Carrano A.V.,

3. Kobayashi A., Olsen A.S., Carrano A.V.,

3. Kobayashi A., Olsen A.S., Carrano A.V.,

4. Kobayashi A., Olsen A.S., Carrano A.V
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Best Local S
Matches 126
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Q9Y4A4;
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Mammalia; Eutheria;
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01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
F22162_1 (Fragment).
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NON_TER
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                                                                              RVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDP----
                                                                                                                         NQALPSGHSKQTQYVLDVQYSPTARIHAS--
                                                                                                                                                             HPAVTG--NLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDAFELTCEAIGKPQPVMVTWV
                                                                                                                                                                                                        LSCLVPRSRPAATLRWYRDRKELKGVSSSQENGKVWSVASTVRFRVDRKDDGGIIICEAQ
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IPR003600; Ig_like.
IPR003006; Ig_MHC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       62;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                 -PTT-
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Matches
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InterPro; IPR003598; I
InterPro; IPR0035000; I
InterPro; IPR003006; I
Pfam; PF00047; ig; 3.
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01-JUN-2001
01-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SMART; SM00409; IG; 3.
SMART; SM00408; IGc2; 3.
SMART; SM00410; IG_like;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fukami T., Maruyama T., Murakami Y.;

'Identification of a murine ortholog of the TSLC1-like gene
submitted (OCT-2001) to the EMBL/GenBank/DDBJ databases.

EMBL; AF195662; AAG35584.1;

EMBL; AV059393; AAL29691.1;

MGD; MGI:2137858; Nec11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zhou Y., Huang X., Yuan "Cloning and expression membrane protein."; Submitted (OCT-1999) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q99N28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunoglobulin SEQUENCE 396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=10090;
[1]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NECL1 OR TSLL1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nectin-like protein 1 (Membrane glycoprotein).
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                                                                                                                               TWVRVDDEMP----QHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPP
                                                                                                                                                                                                                    CQVEHPAVTG-NLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDAFELTCEAIGKPQPVMV
                                                                                                                                                                                                                                                                                                                 EVNCTAMASKPATTIRWFKGNKELKG-KSEVEEWSD--MYTVTSQLMLKVHKEDDGVPVI 181
:||: |||| : |||| : : || : :|| : :||:|| : :||
--- PVPSSSST
                                             TTIPPPTTTT
                                                                                               VWVKEGSEPPLKMTQESALIFP-----FLNKSDSGTYGCTATSNMGSYTAYFTLNVNDPS
                                                                                                                                                                                              CSVNHESLKGADRSTSQRIEVLYTPTAMIR----PEPAHPREGQKLLLHCEGRGNPVPQQY
                                                                                                                                                                                                                                                                                        TLNCQSSGSKPAAQLTWRKGDQELHGDQTRIQEDPNGKTFTVSSSVSFQVTREDDGANIV
                                                                                                                                                                                                                                                                                                                                                                                    TPHELSISISNVALADEGEYTCSIFTMPVRTAKSLVTVLGIPQKPIITGYKSSLREKETA 146
                                                                                                                                                                                                                                                                                                                                                                                                                  SSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVEGEEI 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DDSQPWTSDETVVAGGTVVLKCQVKDHEDSSLQWSNPAQQTLYFGEKRALRDNRIQLVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               109;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity 35.0
09; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     domain.
AA; 42964 MW;
                                             307
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Rodentia;
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Ig_c2.
Ig_like.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27.2%; Score 480.5; 35.0%; Pred. No. 6.86 tive 64; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ig_MHC
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analysis c
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17, Last sequence update)
21, Last annotation updat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cladf8B57D141F3A CRC64;
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of п
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RESULT Q9UJP1

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Best Local Sim
Matches 104;
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SMART; SM00294; 4.1m; 1.
SMART; SM00408; IGc2; 1.
SMART; SM00410; IG_11ke; 2.
Immunoglobblin domain.
SEQUENCE 432 AA; 47020 MW
                                              O9D006 PRELIMINARY;
O9D006;
O1-JUN-2001 (TrEMBLrel. 1
O1-JUN-2001 (TrEMBLrel. 1
O1-DEC-2001 (TrEMBLrel. 1
2610301B19Rik protein.
2610301B19RIK.
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
BK134P22.1 (Novel protein similar to mouse IMMUNOSUPERFAMILY protein
BL2) (NECTLN-11ke protein 1).
BK134P22.1 OR NECL1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q9UJP1;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Zhou Y., Du G., Yuan J., Qlang B.;
Submitted (APR-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AL035403; CAB56227.1; -.
EMBL; AF062733; AAD17540.2; -.
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                        Mus musculus (Mouse)
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InterPro: IPR003600; Ig_Ilke.
InterPro: IPR00306; Ig_MHC.
InterPro: IPR003885; Neurexin-like.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SSGSKPAARLTWRKGDQELHGEPTRIQEDPNGKTFTVSSSVTFQVTREDDGASIVCSVNH
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    Metazoa;
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    Chordata;
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  Craniata;
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RA Arakawa T., Hara A., Fukunishi Y., Konnoo H., Adachi J., Pukuda S.,
RA Alaawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Asito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
RA Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
RA Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Blake J., Boitelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Blake J., Boitt C., Fletcher C., Fujita M., Gariboldi M.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
Lyons P., Marchiomi L., Mashima J., Mazzarelli J., Mombaerts P.,
RA Lyons P., Marchiomi L., Mashima J., Mazzarelli J., Mombaerts P.,
RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,
RA Sasaki H., Sato K., Wang K.H., Weitz C., Whittaker C., Wilming L.,
RA Wynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
RA Wynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
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Best Local S
Matches 93
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SMART; SM00410; IG_like; 1.
SEQUENCE 549 AA; 60703 MW;
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InterPro; IPR003599; Ig
InterPro; IPR003600; Ig_like.
InterPro; IPR003006; Ig_MHC.
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YISDPPTTTTLQPTVQWHSSPADVQDIATEHKKLPFPLSTLATLKDDTIGTIIASVVGG
                                                                                                                                           DANPPPFKSVWSRLDGQWPDGLLASDNTLHFVHPLTVNYSGVYVCKVSNSLGQRSDQKVI
                                                                                                                                                                                                                                                                               TREARGRRITCVVKHPALEKDIRYSFILDIQYAPEVSVTGYDGNWFVGRKG--VNLKCNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DSRFQLLNFSSSELKVSLTNVSISDEGRYFCQLYTDP--PQESYTTITVLVPPRNLMIDI 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GSIIVEPHVTAVWGKNYSLKCLIEVNETITQISWEKIHGKSTQTVAVHHPQYGFSVQGDY
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25.9%;
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Pred. No. 1.7e-16;
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Search completed: November 20, 2002, 07:39:02 Job time: 29.5 secs

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Listing first 45 summaries
1: \SIDS2\gcgdata\geneseq\geneseq\end{1} - \side{1} \text{AA1980.DAT:*}
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Copyright (c) 1943 - 2002 Compug
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

### SUMMARIES

10	9	80	7	σ	Ç	4	w	N	1	Result No.
1694	1741	1741	1741	1741	1741	1741	1741	1741	1765	Score
96.0	98.6	98.6	98.6	98.6	98.6	98.6	98.6	98.6	100.0	Query Match
364	442	442	442	442	440	440	440	402	423	Query Match Length DB
21	23	21	21	21	22	21	20	22	21	B
AAB25586	AAE19887	AAY45092	AAY94341	AAB25619	AAU29040	AAB01321	AAY17830	AAM23691	AAY45093	ID
Protein encoded by	Human tumour suppr	Human lymphoid der	Human cell surface	Protein encoded by	Human PRO polypept	Human PRO355 polyp	Human PRO355 prote	Human EST encoded	Mouse lymphoid der	Description

45	44	43	42	41	40	39	38	37	36	ω 5	34	ω	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11
457	67	67	67	67	67	67	67	69	59	469.5	59	59	469.5	469.5	469.5	469.5	469.5	469.5	469.5	469.5	469.5	469.5	480.5	480.5	78	80	599	606	625.5	25	631	1124.5		1592.5
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AAY53272	AAY94404	ABB50413	AAY45094	AAY69286	AAY94402	ABB51280	ABB51281	AAY94406	AAE00867	ABB84838	ABB95444	ABP61823	AAB53083	AAB80226	AAU12345	AAY45095	AAY69288 .	AAY94403	AAY13358	AAY29592	AAW80405	AAY94405	AAE00868	AAY69287	AAB61142	ABG66677	AAM79402	ABB11897	AAB47251	AAY33741	AAM78418	AAB25593	AAY53028	AAB88427
Human Beat-like 1	Human ACAM6/IgG4-F	Human secreted pro	Human LDCAM bindin	Amino acid sequeun			Human secreted pro	Human ACAM4/IgG4-F	-	-	Human angiogenesis	Human polypeptide	angioge		Human PRO258 polyp	3		ACAM	acid s	Human MBGP1 protei	Ħ	Human ACAM4/IgG1-F		acid s	NOV12	Human novel polype	protein s	Human F22162_1 hom	Beta-secretase. H	Beta-secretase. H	Human protein SEQ		secreted	Human membrane or

# ALIGNMENTS

RESULT 1 AAY45093 Mouse lymphoid derived dendritic cell adhesion molecule. 31-MAY-2000 (first entry) AAY45093; AAY45093 standard; Protein; 423 AA.

Lymphoid derived dendritic cell adhesion molecule; LDCAM; mouse; B7-1; B7-L1; T cell proliferation; natural killer cell; NK; tumour cell; biological activity; quality control reagent; treatment; inflammation; immune system disorder; autolmmune; viral infection; infectious disease; organ transplant rejection; bone marrow; modulator; immune response.

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/note= "N-Glycosylation site"		FT
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/note= "N-Glycosylation site"		FΤ
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9597	Modified-site	FΤ
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8385	Modified-site	FT
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4951	Modified-site	FT
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1356	Domain	FΤ
Location/Qualifiers	Key	FΗ
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organ transplant rejection; bone marrow; modulator; immu	organ trans	X
immune system disorder; autoimmune; viral infection; inf	immune syst	X
activity; quality control reagent; treatment;	biological	ΚW
ell proliferation; natural killer cell; NK; tu	B7-L1; T C	ΧX
Lymphoid derived dendritic cell adhesion molecule; LDCAM	Lymphoid de	ΧX
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Mouse lymphoid derived dendritic cell adhesion molecule.	Mouse lympl	DE
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                                                                                                                                                                                                                                                                                                                                                                                                                                             themselves and B7L-1, an LDCAM binding protein and increases natural killer (NK) cell populations. It may be used to measure the biological activity and as quality control reagents of LDCAM binding proteins. LDCAM may be used for treating disorders associated with malfunctioning of immune system, inflammation, autoimmune disorders, viral infected cells, infectious diseases and for killing tumour cells. They are also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present amino acid sequence is the mouse lymphoid derived dendritic cell adhesion molecule, LDCAM. It is found on lymphoid derived dendritic cells and displays homology to adhesion molecules, B7-1 and cytoplasmic region of B7-L1. Mouse LDCAM is found on whole embryo, testes, triple negative cells murine spienic and lymph node CD8+, $49.1 and dendritic cells. LDCAM polypeptides interacts with T cell surface molecules to alter signalling and inhibits T cell proliferation, bind to
                                                                                                                                                                                                                                                                                                                                                                                            useful for prevention or reducing the effect of organ and bone marrow transplant rejection and for modulating T cell immune responses. LDCAM polypeptides may also be used as carriers for delivering agents attached to T cells or cells bearing B7L-1.
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                                                                                                   ICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDAFELTCEAIGKPQPVMV
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Pred. No. 8.2e-124;
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17-JUL-2000;
03-AUG-2000;
15-SEP-2000;
                                                                                                                                                                                                                             cow, pig, hamster, monkey, macaque, yeast, bacteria, fruit fly, sea urchin and tomato. These were derived from expressed sequence tags from the organism of interest. They can be used in diagnostics, forensics, gene mapping, identification of mutations, to assess biodiversity and for nutritional purposes. The present sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; sheep; pig; cow; fruit fly; yeast; hamster; macaque; tomato; monkey; dog; sea urchin; expressed sequence tag; ESI diagnostics; forensic test; gene mapping; genetic disorder; biodiversity; gene therapy; nutrition.
                                                                                                                                                                                                                     protein of the
                                                                                                                                                                                                                                                                              The present invention provides the protein and coding sequences of novel proteins from a variety of organisms, including human, dog, cat, horse,
                                                                                                                                                                                                                                                                                                             Claim
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antibodies and research use
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         ICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDAFELTCEAIGKPQPVMV
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secreted protein; transmembrane protein; inflammation disorder.
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                                                                                                                                                        identification of homologues, raising antibodies and design of probes and primers. They can be used in a range of diseases related to proteins that they have homology with, e.g. a PRO protein having homology to complement proteins may be used in inflammatory responses.
                                                                                                                                                                                                                                                                                               transmembrane
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                                                                                                                                                                                                                          present invention describes nucleic acids encoding PRO secreted and numembrane proteins used therapeutically. The PRO proteins have ostatic, anti-inflammatory, anti-proliferative and immunosuppressive ivity. The proteins and polynucleotides can be used in therapy, ntification of homologues, raising antibodies and design of probes
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DB; AAX80055.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antibody; screening; detection;
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99..103
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65..69
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                                                                                         "N-myristoylation site"
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"N-myristoylation site"
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inhibition; probe; primer; human.
                                                                                                                                   kinase phosphorylation
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Query Match
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Matches 332
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16-DEC-1998;
22-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                     New human nucleic acids encoding secreted and transmembrane polypeptides which are designated as PRO polypeptides are described The membrane-bound proteins have various industrial applications, including as pharmaceutical and diagnostic agents. The membrane-bound proteins can also be employed for screening of potential peptide or small molecule inhibitors of the relevant receptor/ligand interaction. Anti-PRO antibodies are useful for the affinity purification of PRO
                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 12;
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PPTTTTTTTTTTTTILTIITDSRAGEEGTIGAVDH
                                                                                             LLNFSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVE
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DB; AAA49563.
                                          TWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIP
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javin IJ,
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428..432
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411..417
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A, Godowski PJ, Grima.
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polypeptides, useful as pharmaceutical
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i PJ, Grimaldi CJ, Gurn
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5.4e-122;
nes 3;
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Pan J,
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yg; cat; pig; goat; rabbit; tumour necrosis factor alpha; TNF-alpi
lood; chondrocyte cell; cell proliferation; cell differentiation;
lrenal; lung; breast; prostate; rectum; cervix; liver; genetic di
                                                             KP, Chen J
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2000US-196820P
2000US-198121P
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Watanabe CK,
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                                                                Wood WI, Zhang
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                                                                Godowski PJ,
ng Z;
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                                                                                       Gurney
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RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequences AAU29024-AAU29328 represent PRO polypeptides of the invention. CC The PRO polypeptides and their associated nucleic acids can be used to detect the presence of a tumour in a mammal by comparing the level of CC expression of a PRO polypeptide in a test sample of cells from the animal CC expression in the test sample indicates the presence of a tumour in the CC mammal. Mammals include dogs, cats, cattle, horses, sheep, pigs, goats and rabbits but are preferably human. The polypeptides can be used to Stimulate tumour necrosis factor (TNP) alpha release from human blood, CC when contacted with it. A specific polypeptide can be used to stimulate to the proliferation or differentiation of chondrocyte cells. The PRO proteins can be used to determine the presence of tumours and also susceptibility to tumour development, particularly addrenal, lung, colon, breast, prostate, rectal, cervical, or liver tumours, in mammalian subjects. The oligonucleotide probes specific for the PRO nucleic acids
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Best Local
immune disorder; Addison's disease; rheumatoid arthritis; dermatitis; multiple sclerosis; inflammatory disorder; inflammatory bowel disease; Crohn's disease; nephritis; hyperproliferative disorder; cardiovascular disorder; coronary arteriosclerosis; myocarditis; cance melanoma; lymphoma; wound healing; human.
                                                                                antirheumatic, dermatological; antiproliferative; antiarteriosclerotic;
anticancer; vulnerary; antiviral; antibacterial; antifungal;
                                                                                                                                                                                                                                                    AAB25619
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                                                                                                                                                                                   21-NOV-2000
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332; Conserv
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                                                                                                                                                  encoded
                                                                                                                  protein; immunosuppressant; anti-inflammatory; antiarthritic.
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                                                                                                                                                 human secreted protein gene #11.
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Pred. No. 5.4e
1; Mismatches
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.4e-122;
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TWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIP

ICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDAFELTCEAIGKPQPVMV

ICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDALELTCEAIGKPQPVMV

180 158

LLNFSSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVE 120

LLNFSSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVE

IPTGDGQNLFTKDVTVIEGEVATISCQVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQ

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IPTGDGQNLFTKDVTVIEGEVATISCQVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQ

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CC AAB25576-B25593. The human secreted proteins have various activities of dependent on the tissues in which they are expressed. Examples of the CC activities of the proteins include: immunosuppressant; CC antivities of the proteins include: immunosuppressant; CC antivital; antibacterial; and antifungal activity. The proteins, CC antivital; antibacterial; and antifungal activity. The proteins, CC antivital; antibacterial; and antifungal activity. The proteins, CC polypeptides, agonists and antagonists may be used to treat prevent CC end/or diagnose various disease, disorders and conditions examples of CC which include: immune disorders e.g. Addison's disease, rheumatoid CC arthritis, dermatitis, and multiple sclerosis; inflammatory disorders (C e.g. inflammatory bowel disease, Crohn's disease and nephritis; CC ardiovascular disorders such as paraproteineemias and purpura; CC cardiovascular disorders such as paraproteineemias and myocarditis; CC cancer e.g. melanoma and lymphoma. The proteins and polynucleotide CC sequences may also be used in wound healing and the treatment of CC infectious diseases. The human secreted protein gene #11 and protein sequences are represented in sequences AAA80616 and AAB2586. Sequences CC AAA80677-A80682 represent genes related to the secreted protein gene#11.
Query Match
Best Local S
Matches 332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated nucleic acid molecules encoding human secreted proteins are used for the prevention, amelioration and treatment of autoimmune, inflammatory, hyperproliferative and cardiovascular disorders, cancer wounds, and infectious diseases -
                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to 12 secreted human proteins and to nucleotide sequences encoding them. The polynucleotide sequences in AAA80006-A80623 encode the 12 secreted protein sequences given
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 al Similarity
332; Conser
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Score 1741; DB 21;
Pred. No. 5.4e-122;
1; Mismatches 3;
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                                     Length 442;
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                                                                                                                                                                                                                                                                                                                                                                                     Peptide
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immunosuppressive; antiarteriosclerotic;
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241
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115
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53
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/label= 1
375..394
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67
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159..22
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57..126
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e; antiarteriosclerotic; antibacterial; antiparasit
nootropic; anticonvulsant; cancer; leukaemia;
toid arthritis; asthma; atherosclerosis; akathesia;
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45..442
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antiparasitic;
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PPTTTTTTTTTTTTLTIITDSRAGEEGTIGAVDH

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241 219 181 159 121 99 61 39

TWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIP

300 278 240 218

ICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDALELTCEAIGKPQPVMV ICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDAFELTCEAIGKPQPVMV GEELEVNCTAMASKPATTIRWFKGNTELKGKSEVEEWSDMYTVTSQLMLKVHKEDDGVPV

TWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIP

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LLNFSSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVE

LLNFSSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVE

GEEIEVNCTAMASKPATTIRWFKGNKELKGKSEVEEWSDMYTVTSQLMLKVHKEDDGVPV

180

158

IPTGDGQNLFTKDVTVIEGEVATISCQVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQ

IPTGDGQNLFTKDVTVIEGEVATISCQVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQ

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Query Match Best Local S Matches 332

Local Similarity

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Score 1741; Pred. No. 5 Mismatches

DB 21; 5.4e-122; DB

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12-NOV-1998;
07-DEC-1998;
08-MAR-1999;
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Hillman
                                                                                                    disorders such as rheumatoid arthritis, asthma and atherosclerosis, bacterial and parasitic infections and neuronal disorders such as akathesia, Alzheimer's disease, multiple sclerosis and epilepsy. Polynucleotides encoding HSCRPs may be used as hybridisation probes diagnose these conditions. Anti-HCSRP antibodies may be used as
                                                                                                                                                                             The present sequence is a novel human cell surface receptor protein (HCSRP) designated HCSRP-8. The nucleotide sequence was identified in Incyte Clone 312256 from the cDNA library LUNGNOT02, which was made fr RNA isolated from lung tissue. A number of Incyte Clones were used to assemble the consensus sequence. BLAST analysis showed that the sequence is homologous to immuno-superfamily protein B12 g3779242. HCSRP and it antagonist are useful for preventing or treating disorders associated with decreased or increased expression or activity of HCSRP. Such disorders include cancers such as leukaemia and melanoma, immune
                                                       antagonists, as a targeting or delivery mechanism for bringing pharmaceutical agents into contact with cells or tissues expressing HCSRP and for diagnosis of HCSRP-related disorders. HCSRP and its
                                                                                                                                                                                                                                                                                                                                                                     New human cell surface receptor protein and polynucleotide useful diagnosis, prevention and treatment of cancer, immune disorders, infection and neuronal disorders \,
Sequence
                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 81-82; 97pp; English.
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RESULT 8
   The present amino acid sequence is the human lymphoid derived dendritic cell adhesion molecule, LDCAM. It is found on lymphoid derived dendritic cells and displays homology to adhesion molecules, B7-1 and cytoplasmic region of B7-1. Human LDCAM is expressed in breast, retha, foetal liver, spleen and heart, lung, muscle, placenta, thyroid and lung carcinoma. LDCAM polypeptides interacts with T cell surface molecules to alter signalling and inhibits T cell proliferation, bind to themselves and B71-1, an LDCAM binding protein and increases natural killer (NK) cell populations. It may be used to measure the biological
                                                                                                                                                                                                                              Baum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lymphoid derived dendritic cell adhesion molecule; LDCAM; human; B7-1; B7-11; T cell proliferation; natural killer cell; NK; tumour cell; biological activity; quality control reagent; treatment; inflammation; immune system disorder; autoimmune; viral infection; infectious disease; organ transplant rejection; bone marrow; modulator; immune response.
                                                                                                                          Claim 7; Page 42-43; 44pp; English
                                                                                                                                                    Novel molecules designated LDCAM are capable T cell function -
                                                                                                                                                                                                                                                                               07-AUG-1998;
                                                                                                                                                                                                                                                                                                        05-AUG-1999;
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67..69
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375..3
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RESULT 9
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                                                                                                                                                                                                                                                                                 Human; hepatocellular carcinoma; tumour suppressor lung cancer 1; TSLC1;
liver; lung; pancreatic cancer; cell proliferative disorder; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                          AAE19887 standard; Protein;
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                                                    WPI; 2002-241913/29
                                                                                                                                                            15-AUG-2001;
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                                                                                                                                                                                                                                           Homo sapiens.
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                                                                                                      (UYJO ) UNIV JOHNS HOPKINS SCHOOL
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332; Conser
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Pred. No. 5.4e-122;
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Detecting cell proliferative disorder associated with tumor suppressor lung cancer (TSLC) 1 in subject, comprises contacting proliferating cell of subject with reagent detecting TSLC1 and detecting modification

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Best Local :
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                                                                                                                                 multiple sclerosis; inflammatory disorder; inflammatory bowel Crohn's disease; nephritis; hyperproliferative disorder; cardiovascular disorder; coronary arteriosclerosis; myocarditical cardiovascular disorder; coronary arteriosclerosis; myocardiovascular disorder; myocardiova
                                                                                                                                                                                                                       antirheumatic, dermatological; antiproliferative; antiarteriosclerotic;
anticancer; vulnerary; antiviral; antibacterial; antifungal;
immune disorder; Addison's disease; rheumatoid arthritis; dermatitis;
WO200029435-A1
                                                       Homo sapiens
                                                                                                             melanoma;
                                                                                                                                                                                                                                                                                                            Secreted protein; immunosuppressant; anti-inflammatory; antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                    Protein encoded by human secreted protein gene
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                                                                                                               Lymphoma;
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Pred. No. 5.4e
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5.4e-122;
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                                                                                                                                                                                               disease;
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CC nucleotide sequences encoding them. The polynucleotide sequences given in AAA80666-A80623 encode the 12 secreted protein sequences given in CC AAB25576-B25593. The human secreted proteins have various activities CC aAB25576-B2593. The human secreted proteins have various activities of the tissues in which they are expressed. Examples of the CC activities of the proteins include: immunosuppressant; CC antirinflammatory; antiarthritic; antirheumatic, dermatological; CC antivital; antibacterial; and antifungal activity. The proteins, CC polypeptides, agonists and antagonists may be used to treat prevent CC and/or diagnose various disease, disorders and conditions examples of CC which include: immune disorders e.g. Addison's disease, rheumatoid CC arthritis, dermatitis, and multiple sclerosis; inflammatory disorders c.g. Addison's disease and nephritis; CC ardiovascular disorders such as paraproteinaemias and purpura; CC hyperproliferative disorders used in wound healing and the treatment of CC cancer e.g. melanoma and lymphoma. The proteins and polynucleotide CC sequences may also be used in wound healing and the treatment of CC sequences are represented in sequences AAA80616 and AAB25586. Sequences CC AAA80677-A80682 represent genes related to the secreted protein gene#11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Isolated nucleic acid molecules encoding human secreted proteins are used for the prevention, amelioration and treatment of autoimmune, inflammatory, hyperproliferative and cardiovascular disorders, cancer, wounds, and infectious diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-387742/33.
N-PSDB; AAA80616.
                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HUMA-) HUMAN GENOME SCI INC
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                                                          364
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                                                            AA;
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  96.0%;
99.4%;
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Score 1694; DB 21; Pred. No. 1.4e-118;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    to 12 secreted human proteins and the
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                                                                                  TWVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIP
                                                                                                                                                        ICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDAFELTCEAIGKPQPVMV
                                 PPTTTTTTTTTTTLTIITDSRA
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363
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Qy В Qy В QΥ

Query Match Best Local

Similarity

Length

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Qy Вb QУ

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RESULT 11
AAB88427
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                                                                                                                                                            CC which encode human secretory or membrane proteins represented by AAF93916
CC which encode human secretory or membrane proteins represented by CC AAB88317 - AAB88419. Included in the invention are primers
CC AAB8317 - AAB88419. Included in the invention are primers
CC CAAF93917 - AAF94295 and AAF62232 - AAF62235 which are used to isolate the CC CDNA sequences of the invention. The invention also includes methods for the production of antibodies directed against the proteins, and cDNA CC sequences, which can be used in vaccines. The polynucleotide sequences and the proteins they encode may be used in the prevention, treatment and CC diagnosis of diseases associated with inappropriate secretory protein/membrane protein expression. The nucleic acids and complementary CC sequences may also be used as DNA probes in diagnostic assays (e.g. polymerase chain reactions (PCR)) to detect and quantitate the CC presence of similar nucleic acid sequences in samples. They may also be used to study the expression and function of secretory proteins/membrane CC polypeptides and their role in metabolism. The polypeptides may be used as antigens in the production of antibodies against them and in assays to clentify modulators (agonists and antagonists) of expression and cutivity. The antibodies may also be used as therapeutic agents for detecting the presence of the CC polypeptides in samples (e.g. by enzyme linked inmunosorbant assay also be used as therapeutic astherapeutic astherapeutic aspents for detecting the presence of the arthritis and dispates which may be treated include rheumatoid
                                                                            Query Match
Best Local
                                                             Matches
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11-JAN-2000;
02-MAY-2000;
                                                                                                                                                        arthritie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; secretory protein; membrane protein; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human membrane
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sapiens
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DB; ANF93854.
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                                                             308; Conservative
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2000JP-0183766.
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                                                                           90.2%;
                                                           Score 1592.5; DB 2:
Pred. No. 6.8e-111;
5; Mismatches 23;
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                                                                                          DB 22;
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02-JUN-1998;
22-JUL-1998;
31-JUL-1998;
10-AUG-1998;
11-AUG-1998;
The present invention describes new human secreted proteins which isolated from adult placenta, adult retina, foetal brain, foetal brain, adult brain adult new tissue, adult testes, and adult lymph node cDNA libraries. The hum
                                                                                                                                                             Jacobs K, McCoy JM,
Merberg D, Treacy M
DiBlasio-Smith E, W
                                                                                     New polynucleotides adult placenta, adul
                                                                                                                           N-PSDB;
                                                                                                                                                                                                                                                                                                                                        07-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 haematopoiesis regulation; tissue growth; activin; inhibin; chemotactic; chemokinetic; haemostatic; thrombolytic; receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; secreted protein; nutritional; cytokine;
differentiation; immune stimulating; vaccine; su
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                                                              Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                         tumour inhibition;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  igand; anti-inflammatory; cadherin; tumour invasion suppressor;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              secreted
                                                             71; Page 416-417;
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98US-0095880.
98US-0096068.
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98US-0087645.
98US-0093712.
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Agostino
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tissue growth; activin; inhibin;
                                                                                                  secreted human
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                                                                                      fetal brain,
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                                                                                                                                                                          Collins-Racie L
Steininger RJ,
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fetal -
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                       foetal kidney,
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Best Local :
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                                                                                                                                                                                                                                                                                     Crohn's disease; nephritis; hyperproliferative disorder; cardiovascular disorder; coronary arteriosclerosis; myoc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (e.g. as vaccines) or suppressing activity, haematopoiesis regulating activity, tissue growth activity, activin/inhibin activity, chemotactic/Chemokhnetic activity, haemostatic and thrombolytic activity, receptor/ligand activity, anti-inflammatory activity, activity, cadherin/tumour invasion suppressor activity, and tumour inhibition activity. The polynucleotides are also stated to be useful for gene therapy. AAZ33316 to AAZ33373 encode human secreted proteins, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        secreted proteins, and the polynucleotides encoding them, are predicted to have biological activities which would make them suitable for treating, preventing or ameliorating medical conditions in humans and animals. Suggested activities include nutritional activity, cytokine and cell proliferation/differentiation activity, immune stimulating
                           27-OCT-1999;
                                                                                 25-MAY-2000
                                                                                                                                          WO200029435-A1
                                                                                                                                                                                                      Homo
                                                                                                                                                                                                                                                                                                                                              multiple sclerosis; inflammatory disorder; inflammatory bowel disease;
                                                                                                                                                                                                                                                                                                                                                                                                       antirheumatic, dermatological; antiproliferative; antiarteriosclerotic;
anticancer; vulnerary; antiviral; antibacterial; antifungal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Protein encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB25593 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                therapy. AAZ33316 to AAY52998 to AAY53060
                                                                                                                                                                                                                                                                                                                                                                               lmmune disorder; Addison's disease; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     332
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LLNFSSSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVE
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                                                                                                                                                                                                                                                            lymphoma;
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                        99WO-US25031.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunosuppressant; anti-inflammatory; antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         human secreted protein gene #11 clone HOUDJ81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Protein;
                                                                                                                                                                                                                                                               wound
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DSRAGEEGSIRAVDH
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Pred. No. 2.5e
1; Mismatches
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.5e-109;
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                                                                                                                                                                                                                                                                                                                                                                               dermatitis;
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and/or diagnose various disease, disorders and conditions examples of which include: immune disorders e.g. Addison's disease, rheumatoid arthritis, dermatitis, and multiple sclerosis; inflammatory disorders e.g. inflammatory bowel disease, Crohn's disease and nephritis; hyperproliferative disorders such as paraproteinaemias and purpura; cardiovascular disorders e.g. coronary arteriosclerosis and myocarditis; cancer e.g. melanoma and lymphoma. The proteins and polynucleotide sequences may also be used in wound healing and the treatment of infectious diseases. The human secreted protein gene #11 and protein sequences are represented in sequences AAA80616 and AAB25586. Sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antiproliferative; antiarteriosclerotic; anticancer; vulnerary; antiviral; antibacterial; and antifungal activity. The proteins, polypeptides, agonists and antagonists may be used to treat prevent
                                                                                                                                         06-NOV-2001
                                                                                                                                                                                                                                                      AAM78418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           activities of the proteins include: immunosuppres anti-inflammatory; antiarthritic; antirheumatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to 12 secreted human proteins and the nucleotide sequences encoding them. The polynucleotide sequences given in AAA80606-A80623 encode the 12 secreted protein sequences given in AAB25576-B25593. The human secreted proteins have various activities
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 685-686; 803pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Isolated nucleic acid molecules encoding human secreted proteins used for the prevention, amelioration and treatment of autoimmuna inflammatory, hyperproliferative and cardiovascular disorders, continuations, in the continuation of the continuati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA80677-A80682 represent genes related to the secreted protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               dependent on the tissues in which they are expressed.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Greene
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95.1%;
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No. 2.86
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vaccine; peptide Human; cytokine;

cell proliferation; cell differentiation; gene th therapy; stem cell growth factor; haematopoiesis;

therapy;

Human protein

SEQ ID NO 1080.

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27-APR-2000; 2000US-0598075.
20-JUN-2000; 2000US-0598075.
19-JUL-2000; 2000US-0620325.
01-SEP-2000; 2000US-0654936.
15-SEP-2000; 2000US-0663561.
20-OCT-2000; 2000US-0693325.
30-NOV-2000; 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to polynucleotides (AAK51456-AAK53435) and the encoded polypeptides (AAM78323-AAM80302) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polynucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopoissis regulating activity, tissue growth factor activity, immunomodulatory activity and activity in activity and may be useful in the diagnosis and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tang YT,
Zhao QA,
Xue AJ,
                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 20; Page 3307-3308; 6221pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acids encoding polypeptides with cytokine-like activities useful in diagnosis and gene therapy -
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                                                                                                                                                                                                                                                                                                                                                                                                                               inflammation.
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DB; AAK51551.
                                                                                                                       IEVNCTAMASKPATTIRWFKGNKELKGKSEVEEWSDMYTVTSQLMLKVHKEDDGVPVICQ 183
            WVRVDDEMPQHAVLSGPNLFINNLNKTDNGTYRCEASNIVGKAHSDYMLYVYDPPTTIPP
                                                                                                                                                                                                                              GAGQEVQTENVTVAEGGVAEITCRLHQYDGSIVVIQNPARQTLFFNGTRALKDERFQLEE 80
                                                                                                                                                                                                                                                                                                                                                                                  missing at the
                                           AQNQALPSGHSKQTQYVLDVQYSPTARIHAS---QAVVREGDTLVLTCAVTGNPRPNQIR
                                                                       VEHPAVTG--NLOTORYLEVOYKPOVHIOMTYPLOGLTREGDAFELTCEAIGKPOPVMVT
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Wang D, Wang J, Z
Yang Y, Wejhrman T,
                                                                                                                                                                                                                                                                                                                                                                                                                                               of cancer, leukaemia, nervous system disorders, arthritis and
                                                                                                                                                                                                                                                                                                                                                       387
                                                                                                                                                                                                                                                                                          Conservative
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m disorder; arthritis; inflammation.
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, Zhang J, Ren F, C
n T, Goodrich R;
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Pred. No. 4e-39;
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Wang ZW;
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RESULT 15
AAY33741
                                             Query Match
           Matches
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07-JUN-1995;
07-JUN-1995;
                                                                                                                  This sequence is the human beta-secretase enzyme polynucleotide. Beta-secretase is capable of cleaving the beta-amyloid precursor protein (APP) (ANY33742;swedish mutant APP). This enzyme is used in a method for detecting human beta-secretase cleavage of polypeptides and for identifying beta-secretase inhibitors. The protein has a deduced molecular weight of 49 kD, although this is without glycosylation and the expected molecular weight is therefore higher. Inhibition of beta-secretase activity would be useful for chemical modelling of a critical event in the pathology of Alzheimer's disease. Inhibitors of beta-secretase would be useful for the prevention and treatment of Alzheimer's disease and Down's Syndrome.
                                                                                                                                                                                                                                                                                                                                                                                          A method for detecting human beta-secretase cleavage of polypeptides useful for identifying beta-secretase inhibitors
                                                                                     Sequence
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GQFPLTQNVTVVEGGTAILTCRVDQNDNTSLQWSNPAQQTLYFDDKKALRDNRIELVRAS GONLETKDVTVIEGEVATISCOVNKSDDSVIQLLNPNRQTIYFRDFRPLKDSRFQLLNFS

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                                66 SSELKVSLTNVSISDEGRYFCQLYTDPPQESYTTITVLVPPRNLMIDIQKDTAVEGEEIE 125
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93 WHELSISVSDVSLSDEGQYTCSLFTMPVKTSKAYLTVLGVPEKPQISGFSSPVMEGDLMQ 152
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Search completed: November 20, 2002, 07:37:57 Job time: 63.5 secs

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Gapop 10.0 , Gapext 0.5
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed. and is derived by analysis of the total score distribution.

## SUMMARIES

19	18	17	16	15	14	13	12	11	10	9	8	7	თ	v	4	w	N		Result No.
1741	1741	1741	1741	1741	1741	1741	1741	1741	1741	1741	1741	1741	1741	1741	1741	1741	1765	1765	Score
98.6	98.6	98.6	98.6	98.6	98.6	98.6	98.6	98.6	98.6	98.6	98.6	98.6	98.6	98.6	98.6	98.6	100.0	100.0	Query Match
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US-09-778-510-20	US-10-052-586-34	US-09-943-851A-61	US-09-944-654-61	US-09-943-762-61	US-09-944-432-61	US-09-944-097-61	US-09-944-396-61	US-09-945-015-61	US-09-945-587-61	US-09-944-862-61	US-09-944-457-61	US-09-944-449-61	US-09-866-028-61	US-09-944-896-61	US-09-944-403-61	US-09-944-413-61	US-09-778-187B-4	US-09-778-510-22	ID
_ `	Sequence 34, Appl	Sequence 61, Appl	Sequence 61, Appl	Sequence 61, Appl	•	•	Sequence 61, Appl	•	Sequence 61, Appl	•	•	Sequence 61, Appl	Sequence 4, Appli	Sequence 22, Appl	Description				

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	100	11.0	11.2	11.5	11.6	11.6	11.7	11.8	11.8	11.8	11.9	12.5	12.5	12.5	12.5	26.5	26.5	26.6	26.6	26.6	26.6	26.6	26.6	27.2	27.2	98.6
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	115-09-808-603-67	US-09-736-457-334	US-09-808-602-71	US-09-925-301-1133	US-09-756-551A-17	US-10-052-586-440	US-09-764-853-605	US-09-764-853-794	US-10-047-542-80	US-10-047-542-90	US-09-919-172-20	US-09-978-697-523	US-09-966-546-6	US-09-966-546-4	US-09-978-295A-523	US-09-778-187B-8	US-09-778-510-2	US-09-909-088B-84	US-09-909-320-84	US-09-745-763-102	US-09-778-187B-10	US-09-778-510-6	US-09-905-291A-84	US-09-778-510-4	US-10-047-542-79	US-09-778-187B-2
neducance of other		Sequence 334, App	Sequence 71, Appl	Sequence 1133, Ap	Sequence 17, Appl	Sequence 440, App	Sequence 605, App	Sequence 794, App	Sequence 80, Appl	Sequence 90, Appl	Sequence 20, Appl	Sequence 523, App	Sequence 6, Appli	Sequence 4, Appli	Sequence 523, App	Sequence 8, Appli	Sequence 2, Appli	Sequence 84, Appl	Sequence 84, Appl	Sequence 102, App	Sequence 10, Appl	6	Sequence 84, Appl	Sequence 4, Appl1	79,	Sequence 2, Appli

## ALIGNMENTS

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; SOFTWARE: PatentIn Ve
; SEQ ID NO 22
; LENGTH: 423
; TYPE: PRT
; ORGANISM: Mus musculus
US-09-778-510-22
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US-09-778-510-22
Sequence 22, Application US/09778510
Patent No. US20020164686A1
Patent No. US20020164686A1
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APPLICANT:
Baum,
TITLE OF INVENTION:
FILE REFERENCE:
                                                                                                                                                                                                                              Query Match 100.0%; Score 1765; DB 9; Best Local Similarity 100.0%; Pred. No. 9.8e-108; Matches 336; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/778,510
CURRENT FILING DATE: 2001-02-07
PRIOR APPLICATION NUMBER: PCT/US99/179
PRIOR FILING DATE: 1999-08-07
                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 19
NUMBER OF SEQ ID NOS:
141
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PatentIn Ver. 2.0
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Gaps

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Sequence 61, Application US/
Patent No. US20020156004A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin
APPLICANT: Botscein, David
APPLICANT: Eaton, Dan
APPLICANT: Filvaroff, Ellen
APPLICANT: Gerritsen, Mary
APPLICANT: Goddard, Audrey
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US-09-778-187B-4
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US-09-778-1878-4
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Best Local Similarity 100.0%; Pred. No. 9
Matches 336; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 4, Application US/09778187B Patent No. US20020168712A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Baum, Peter R.
APPLICANT: Fanslow III, Willi
TITLE OF INVENTION: MOLECULES
FILE REFERENCE: 2873-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: PCT/US99/17905
PRIOR FILING DATE: 1999-08-05
PRIOR APPLICATION NUMBER: US 60/095,672
PRIOR FILING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/778,187B CURRENT FILING DATE: 2001-02-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.1
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                                        Filvaroff, Ellen
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DESIGNATED LDCAM
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PRIOR APPLICATION NUMBER: 09/866,028
PRIOR FILING DATE: 2001-05-25
PRIOR PPLICATION NUMBER: 60/067,411
PRIOR PFLING DATE: December 3, 1997
PRIOR PFLING DATE: December 11, 1997
PRIOR PFLING DATE: December 11, 1997
PRIOR PFLING DATE: December 11, 1997
PRIOR FILING DATE: December 12, 1997
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CURRENT APPLICATION NUMBE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: WOOD, WILLIAM
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE
TITLE OF INVENTION: ACIDS ENCODING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR
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                                                                                                                                                                                                                                                                                                                                                                                                                     OR FILING DATE: December 18, 1997
OR APPLICATION NUMBER: 60,0070,440
OR FILING DATE: January 5, 1998
OR APPLICATION NUMBER: 60,074,086
OR FILING DATE: February 9, 1998
OR APPLICATION NUMBER: 60,074,092
OR FILING DATE: February 9, 1998
OR APPLICATION NUMBER: 60,075,945
OR FILING DATE: February 25, 1998
OR APPLICATION NUMBER: 60,075,945
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 60,7112,850
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 60,714,296
OR APPLICATION NUMBER: 60,7146,222
OR FILING DATE: July 28, 1999
OR APPLICATION NUMBER: 60,7146,222
OR FILING DATE: July 28, 1999
OR APPLICATION NUMBER: 60,7146,222
OR FILING DATE: July 28, 1999
OR FILING DATE: July 28, 1999
OR APPLICATION NUMBER: 60,7146,222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DR FILING DATE: December 16, 1997
DR APPLICATION NUMBER: 60/069,702
DR FILING DATE: December 16, 1997
DR APPLICATION NUMBER: 60/069,870
DR FILING DATE: December 17, 1997
DR APPLICATION NUMBER: 60/069,873
DR FILING DATE: December 17, 1997
DR APPLICATION NUMBER: 60/069,873
DR FILING DATE: December 17, 1997
                                                                                                                                                                       APPLICATION NUMBER: 09/254,311
APPLICATION NUMBER: 09/254,311
FILING DATE: March 3, 1999
A APPLICATION NUMBER: PCT/US99/12252
APPLICATION NUMBER: PCT/US99/12252
                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 60/146,222
FILLING DATE: July 28, 1999
APPLICATION NUMBER: PCT/US98/19330
FILING DATE: September 16, 1998
APPLICATION NUMBER: PCT/US98/25108
                      APPLICATION NUMBER: PCT/US99/28301 FILING DATE: December1, 1999
                                                             APPLICATION NUMBER: PCT/US99/28313 FILING DATE: No. US20020156004Alember
                                                                                                    APPLICATION NUMBER: PCT/US99/28409 FILING DATE: No. US20020156004Alember
                                                                                                                                           APPLICATION NUMBER: PCT/US99/21090 FILING DATE: September 15, 1999
                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 09/216,021 FILING DATE: December 16, 1998 APPLICATION NUMBER: 09/218,517
                                                                                                                                                                                                                                                                                                                                               FILING DATE:
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FILING DATE: December 16, 1997
APPLICATION NUMBER: 60/069,694
FILING DATE: December 16, 1997
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Hillan, Kenneth
Kljavin, Ivar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tumas,Daniel
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RESULT 4
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; ORGANISM: Homo
US-09-944-413-61
                                                           GENERAL INFO
APPLICANT:
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Best Local Similarity
Matches 332; Conserv
                                                                                                                                                                                                                                    Sequence 61, Application US/09944403
Patent No. US20020165143A1
GENERAL INFORMATION:
 APPLICANT:
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OR APPLICATION NUMBER: PCT/US00/04414
OR FILING DATE: February 22, 2000
OR APPLICATION NUMBER: PCT/US00/05841
OR FILING DATE: March 2, 2000
OR APPLICATION NUMBER: PCT/US00/08439
OR FILING DATE: March 30, 2000
OR APPLICATION NUMBER: PCT/US00/14042
OR FILING DATE: March 30, 2000
OR APPLICATION NUMBER: PCT/US00/14042
OR APPLICATION NUMBER: PCT/US00/14042
OR APPLICATION NUMBER: PCT/US00/20710
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Goddard, Audrey
Godowski, Paul
Grimaldi, Christopher
Roy, Margaret
Tumas, Daniel
                                                                           Gurney, Austin
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                                               Kljavin, Ivar
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E: December 16, 1998
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E: December 16, 1997
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Best Local Similarity
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P2548B1C1 CURRENT APPLICATION NUMBER: US/09/944,896 CURRENT FILING DATE: 2001-08-31 PRIOR APPLICATION NUMBER: 09/866,028 PRIOR FILING DATE: 2001-05-25
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PR APPLICATION NUMBER: PCT/US00/14042

DR FILING DATE: May 22, 2000

DR APPLICATION NUMBER: PCT/US00/20710

DR FILING DATE: July 28, 2000

DR APPLICATION NUMBER: PCT/US00/32678

DR FILING DATE: December 1, 2000

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Grimaldi, Christopher
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Goddard, Audrey
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DR APPLICATION NUMBER: 09/216,021
DR FILING DATE: December 16, 1998
DR APPLICATION NUMBER: 09/218,517
DR FILING DATE: December 22, 1998
DR APPLICATION NUMBER: 09/254,311
DR FILING DATE: March 3, 1999
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DR APPLICATION NUMBER: 60/074,086
DR FILLING DATE: February 9, 1998
DR APPLICATION NUMBER: 60/074,092
DR FILLING DATE: February 9, 1998
DR APPLICATION NUMBER: 60/075,945
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                                               APPLICATION NUMBER: PCT/US00/32678 FILING DATE: December 1, 2000
                                                                                                                         FILING DATE: March 30, 2000 APPLICATION NUMBER: PCT/US00/14042
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APPLICATION NUMBER: PCT/US99/28409
FILING DATE: No. US20020168715Alember
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E: December 16, 1997
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NUMBER: PCT/US00/20710
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APPLICANT: Baker, Ke
APPLICANT: Betstein
APPLICANT: Eaton, Da
APPLICANT: Ferrara,
APPLICANT: Ferrara,
APPLICANT: Gerritse
APPLICANT: Goddard,
APPLICANT: Goddwski
APPLICANT: Grimaldi
APPLICANT: Grimaldi
APPLICANT: Hillan,
APPLICANT: Napler, M
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Best Local Similarity
Matches 332; Conserv
 Best Local Similarity Matches 332; Conserva
                       Query Match
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APPLICANT: Tumas Daniel
APPLICANT: Tumas Daniel
APPLICANT: Wood, Milliam
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P2548P1C1
CURRENT APPLICATION NUMBER: US/09/866,028
CURRENT FILING DATE: 2001-05-25
                                                              LENGTH: 440
TYPE: PRT
ORGANISM: Homo 9
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           Score 1741; DB 10;
Pred. No. 3.7e-106;
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Pred. No. 3.7e-106;
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P2548PIC1
CURRENT APPLICATION NUMBER: U$/09/944,449
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 09/866,028
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December 3, 1997
PRIOR APPLICATION NUMBER: 60/069,335
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,327
PRIOR APPLICATION NUMBER: 60/069,278
PRIOR APPLICATION NUMBER: 60/069,278
PRIOR APPLICATION NUMBER: 60/069,278
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,696
PRIOR FILING DATE: December 16, 1997
PRIOR APPLICATION NUMBER: 60/069,696
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PRIOR APPLICATION NUMBER: 60/069,696
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PRIOR FILING DATE: December 16, 1997
PRIOR APPLICATION NUMBER: 60/069,890
PRIOR FILING DATE: December 16, 1997
PRIOR APPLICATION NUMBER: 60/069,890
PRIOR FILING DATE: December 17, 1997
PRIOR APPLICATION NUMBER: 60/069,870
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APPLICANT: Baker, F
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Roy, Margaret
Tumas, Daniel
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Godowski, Paul
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APPLICATION NUMBER: 60/068,017 FILING DATE: December 18, 1997 APPLICATION NUMBER: 60/070,440 FILING DATE: January 5, 1998 APPLICATION NUMBER: 60/074,086

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PRIOR APPLICATION NUMBER: 09/218,517
PRIOR APPLICATION NUMBER: 09/218,517
PRIOR APPLICATION NUMBER: 09/254,311
PRIOR APPLICATION NUMBER: 09/254,311
PRIOR FILING DATE: March 3, 1999
PRIOR APPLICATION NUMBER: DET/US99/12252
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IOR FILING DATE: September 15, 1999
IOR FILING DATE: September 15, 1999
IOR APPLICATION NUMBER: PCT/US99/28409
IOR FILING DATE: No. US200/2010/2647Alember 30, 19
IOR APPLICATION NUMBER: PCT/US99/28313
IOR FILING DATE: No. US200/2010/2647Alember 30, 19
IOR APPLICATION NUMBER: PCT/US99/28301
IOR APPLICATION NUMBER: PCT/US99/28301
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                                                             APPLICATION NUMBER: 60/074,092
FILING DATE: February 9, 1998
FILING DATE: February 0
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APPLICATION NUMBER: 09/216,021
FILING DATE: December 16, 1998
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APPLICATION 1
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PRIOR APPLICATION NUMBER: 60/067,4
PRIOR APPLICATION NUMBER: 60/067,4
PRIOR ETLING DATE: December 3, 199
PRIOR APPLICATION NUMBER: 60/069,4
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CURRENT FILING DATE: 2001-09-26
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P2548PIC1
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OR FILING DATE: December 12, 1997
OR APPLICATION NUMBER: 60/069,696
OR FILING DATE: December 16, 1997
OR APPLICATION NUMBER: 60/069,694
OR FILING DATE: December 16, 1997
OR APPLICATION NUMBER: 60/069,702
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FILING DATE: December 11, 1997
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APPLICATION FILING DATE:
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Gurney,Austin
Hillan,Kenneth
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                                                                                                                                          NUMBER: 60/069, 19 NUMBER: 60/069, 8
               : February 9, 1:
NUMBER: 60/074
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NUMBER: 60/070,
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TYPE: PRT
ORGANISM: Homo Sapien
1S-09-944-457-61
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Best Local Similarity 98.8
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IOR APPLICATION NUMBER: 09/216,021
IOR FILING DATE: December 16, 1998
IOR FILING DATE: December 22, 1998
IOR APPLICATION NUMBER: 09/254,311
IOR FILING DATE: December 22, 1998
IOR APPLICATION NUMBER: 09/254,311
IOR FILING DATE: March 3, 1999
IOR FILING DATE: March 3, 1999
IOR FILING DATE: June 22, 1999
IOR APPLICATION NUMBER: PCT/US99/21090
IOR FILING DATE: September 15, 1999
IOR APPLICATION NUMBER: PCT/US99/28409
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APPLICATION NUMBER: PCT/US00/05841
FILING DATE: March 2, 2000
APPLICATION NUMBER: PCT/US00/08439
FILING DATE: March 30, 2000
APPLICATION NUMBER: PCT/US00/14042
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APPLICATION NUMBER: PCT/US00/03565
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APPLICATION NUMBER: PCT/US00/04414
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                 ICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDAFELTCEAIGKPQPVMV 240
                                                                      GEEIEVNCTAMASKPATTIRWFKGNTELKGKSEVEEWSDMYTVTSQLMLKVHKEDDGVPV
ICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDALELTCEAIGKPQPVMV
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Pred. No. 3.7e-106;
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APPLICANT: Baker,
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APPLICANT: Eaton,
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PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,411
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CURRENT FILING DATE: 2001-09-26
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APPLICATION NUMBER: 60,074,094

OR APPLICATION NUMBER: 60,075,945

"TILING DATE: February 9, 1998

"TILING DATE: February 9, 1998

"TILING DATE: February 5, 1998

"ATTEL February 25, 1998

"MUMBER: 60,112,850

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OR FILING DATE: December 16, 1997
OR APPLICATION NUMBER: 60/069,702
OR FILING DATE: December 16, 1997
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APPLICATION NUMBER: 60/069335
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APPLICATION NUMBER: 60/069,278
FILING DATE: December 11, 1997
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APPLICATION NUMBER: 60/068,
FILING DATE: December 18, 1
APPLICATION NUMBER: 60/070,
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Gurney,Austin
Hillan,Kenneth
Kljavin,Ivar
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Godowski, Paul
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Roy, Margaret
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NUMBER: 6
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FILING DATE:
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FILING DATE:
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NUMBER: PCT/US99/28301
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PRIOR APPLICATION NUMBER: 09/866,028
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,335
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APPLICANT: Botste,
APPLICANT: Eaton, I
APPLICANT: Ferrarr
APPLICANT: Filvarr
APPLICANT: Gerrit:
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P2548P1C1
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                                                                                                                                            OR FILING DATE: December 18, 1997
OR APPLICATION NUMBER: 60/070,440
OR FILING DATE: January 5, 1998
OR APPLICATION NUMBER: 60/074,086
OR FILING DATE: February 9, 1998
OR APPLICATION NUMBER: 60/074,096
OR FILING DATE: February 9, 1998
OR APPLICATION NUMBER: 60/075,945
OR FILING DATE: February 25, 1998
OR APPLICATION NUMBER: 60/112,945
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 60/113,296
OR APPLICATION NUMBER: 60/146,222
OR FILING DATE: December 22, 1998
OR APPLICATION NUMBER: 60/146,222
OR FILING DATE: JULY 28, 1999
OR APPLICATION NUMBER: 60/146,222
OR FILING DATE: JULY 28, 1999
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APPLICATION NUMBER: 60/069,873
FILING DATE: December 17, 1997
APPLICATION NUMBER: 60/068,017
  FILING DATE: APPLICATION |
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APPLICATION NUMBER: 60/069,870
FILING DATE: December 17, 1997
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FILING DATE: December 11, 1997
APPLICATION NUMBER: 60/069,425
FILING DATE: December 12, 1997
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RESULT 11
US-09-945-015-61
Sequence 61, Application US/09945015
Patent No. US20020132768A1
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US-09-945-587-61
                                                            GENERAL INFORMATION:
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OR APPLICATION NUMBER: PCT/US00/05841
OR FILING DATE: March 2, 2000
OR APPLICATION NUMBER: PCT/US00/08439
OR FILING DATE: March 30, 2000
OR APPLICATION NUMBER: PCT/US00/14042
OR FILING DATE: May 22, 2000
OR APPLICATION NUMBER: PCT/US00/20710
OR APPLICATION NUMBER: PCT/US00/20710
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OR FILING DATE: No. US20020127643A1ember 30
OR APPLICATION NUMBER: PCT/US99/28301
OR FILING DATE: December1, 1999
OR APPLICATION NUMBER: PCT/US99/30095
OR FILING DATE: December 16, 1999
OR PILING DATE: December 17, 1999
OR PILING DATE: December 17, 2000
OR APPLICATION NUMBER: PCT/US00/04414
OR APPLICATION NUMBER: PCT/US00/04414
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FILING DATE: July 28,
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FILING DATE: NO. US20020127643Alember
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FILING DATE: March 3,
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Baker, Kevin
Botstein, David
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3, 1999
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Pred. No. 3.7e-106;
1; Mismatches 3;
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 09/966,028
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR APPLICATION NUMBER: 60/069,335
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PRIOR APPLICATION NUMBER: 60/069,278
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,425
PRIOR FILING DATE: December 12, 1997
PRIOR APPLICATION NUMBER: 60/069,696
PRIOR FILING DATE: December 16, 1997
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                                           OR APPLICATION NUMBER: 09/254,311
OR APPLICATION NUMBER: 09/254,311
IOR FILING DATE: MATCh 3, 1999
AIOR APPLICATION NUMBER: PCT/US99/12252
ATTING DATE: June 22, 1999
TTING DATE: MIMBER: PCT/US99/21090
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Kljavin, Ivar
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Gerritsen,Mary
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NUMBER: 60/074, C
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NUMBER: PCT
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NUMBER: PCT/US98/25108
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NUMBER: 60/112, 850

: December 16, 1998

NUMBER: 60/113, 296

: December 22, 1998

NUMBER: 60/146, 222
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Sequence 61, Application US/09944396
Patent No. US20020132981A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin
APPLICANT: Botstein, David
APPLICANT: Eaton, Dan
APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Gerritsen, Mary
APPLICANT: Gerritsen, Mary
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Paul
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US-09-944-396-61
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PRIOR FILING DATE: March 2, 2000
PRIOR FILING DATE: March 30, 2000
PRIOR FILING DATE: March 30, 2000
PRIOR FILING DATE: May 22, 2000
PRIOR PRIOR DATE: May 22, 2000
PRIOR APPLICATION NUMBER: PCT/US00/14042
PRIOR FILING DATE: July 28, 2000
PRIOR APPLICATION NUMBER: PCT/US00/20710
PRIOR FILING DATE: July 28, 2000
PRIOR APPLICATION NUMBER: PCT/US00/32678
PRIOR FILING DATE: December 1, 2000
PRIOR APPLICATION NUMBER: PCT/US01/06520
PRIOR APPLICATION NUMBER: PCT/US01/06520
PRIOR FILING DATE: February 28, 2001
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OR FILING DATE: NO. US20020133768Alember 30, 19:
OR APPLICATION NUMBER: PCT/US99/28301
OR FILING DATE: December1, 1999
OR APPLICATION NUMBER: PCT/US99/30095
OR FILING DATE: December 16, 1999
OR APPLICATION NUMBER: PCT/US90/03565
OR FILING DATE: Pebruary 11, 2000
OR APPLICATION NUMBER: PCT/US00/04414
OR APPLICATION NUMBER: PCT/US00/04414
OR FILING DATE: Pebruary 22, 2000
OR FILING DATE: Pebruary 22, 2000
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Godowski,Paul
Grimaldi,Christopher
Gurney,Austin
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PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
PRIOR FILING DATE: December 11, 1997
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069,335
PRIOR FILING DATE: December 11, 1997
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: p.5548plc1
CURRENT APPLICATION NUMBER: US/09/944,396
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CURRENT FILING DATE: 2001-09-26
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DR APPLICATION NUMBER: 09/218,517
DR FILING DATE: December 22, 1998
DR APPLICATION NUMBER: 09/254,311
DR FILING DATE: MATCh 3, 1999
DR APPLICATION NUMBER: PCT/US99/12252
DR FILING DATE: June 22, 1999
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OR FILING DATE: December 16, 199
OR APPLICATION NUMBER: 60/069,879
OR FILING DATE: December 17, 1997
OR FILING DATE: December 17, 1997
OR APPLICATION NUMBER: 60/069,873
OR FILING DATE: December 18, 1997
OR APPLICATION NUMBER: 60/070,440
OR FILING DATE: December 18, 1997
OR APPLICATION NUMBER: 60/070,440
OR FILING DATE: January 5, 1998
OR APPLICATION NUMBER: 60/074,086
OR FILING DATE: February 9, 1998
OR APPLICATION NUMBER: 60/074,092
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APPLICATION NUMBER: 60/112,850
FILING DATE: December 16, 1998
APPLICATION NUMBER: 60/113,296
FILING DATE: December 22, 1998
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Kljavin, Ivar
                     NUMBER:
                                      December 16,
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Best Local Similarity
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FILING DATE: December 1, 2000
APPLICATION NUMBER: PCT/US01/06520
FILING DATE: February 28, 2001
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FILING DATE: February 22, 2000
APPLICATION NUMBER: PCT/US00/05841
FILING DATE: March 2, 2000
APPLICATION NUMBER: PCT/US00/08439
FILING DATE: March 30, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GEEIEVNCTAMASKPATTIRWFKGNKELKGKSEVEEWSDMYTVTSQLMLKVHKEDDGVPV 180
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                                                                                                                                                                                                                                                                 INFORMATION
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 INVENTION:
                         Napier, Mary
Roy, Margaret
Tumas, Daniel
Wood, William
                                                                                                                Gurney, Austin
                                                                                                                                Grimaldi, Christopher
                                                                                                                                            Godowsk1, Paul
                                                                                                                                                           Goddard, Audrey
                                                                                                                                                                          Gerritsen, Mary
                                                                                                                                                                                       Filvaroff, Ellen
                                                                                     Kljavin, Ivar
                                                                                                    Hillan, Kenneth
                                                                                                                                                                                                       Ferrara, Napoleone
                                                                                                                                                                                                                        Eaton, Dan
                                                                                                                                                                                                                                    Botstein, David
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SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC ACIDS ENCODING THE SAME
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Pred. No. 3.7e-106;
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PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069335
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CURRENT FILING DATE: 2001-08-31
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         OR FILING DATE: February 2
DR APPLICATION NUMBER: PCT
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APPLICATION NUMBER: 60/070,4
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NUMBER: PCT/US99/12252
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NUMBER: PCT/US98/25108
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NUMBER: 60/112,
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PRIOR APPLICATION NUMBER: 09/866,028
PRIOR FILING DATE: 2001-09-26
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December 11, 1997
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PRIOR APPLICATION NUMBER: 60/069336
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APPLICANT: Baker, Ke
APPLICANT: Botstein
APPLICANT: Eaton, Da
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PRIOR FILING DATE: December 1, 2000
PRIOR APPLICATION NUMBER: PCT/US01/06520
PRIOR FILING DATE: February 28, 2001
NUMBER OF SEQ ID NOS: 120
SEQ ID NO 61
LENGTH: 440
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                                                                                                                                                     APPLICANT: WOOD, WILLIAM
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P2548P1C1
CURRENT APPLICATION NUMBER: US/09/944,432
CURRENT FILING DATE: 2001-09-26
CURRENT FILING DATE: 2001-09-26
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Grimaldi,Christopher
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Tumas, Daniel
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Pred. No. 3.7e-106;
1; Mismatches 3;
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SEQ ID NO 61
LENGTH: 440
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OR APPLICATION NUMBER: 60/074,086

FILING DATE: February 9, 1998

FOR FILING DATE: February 9, 1998

FOR FILING DATE: February 9, 1998

FOR FILING DATE: February 25, 1998

FOR FILING DATE: February 25, 1998

FOR FILING DATE: December 16, 1998

FOR APPLICATION NUMBER: 60/112,850

FOR FILING DATE: December 22, 1998

FOR APPLICATION NUMBER: 60/146,222

FOR FILING DATE: December 22, 1998

FOR APPLICATION NUMBER: 60/146,222

FOR FILING DATE: December 16, 1998

FOR APPLICATION NUMBER: PCT/US98/25108

FOR APPLICATION NUMBER: PCT/US98/25108

FOR FILING DATE: September 1, 1998

FOR APPLICATION NUMBER: PCT/US98/25108

FOR FILING DATE: December 1, 1998

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OR APPLICATION NUMBER: 60/069,278
OR FILING DATE: December 11, 1997
OR APPLICATION NUMBER: 60/069,425
OR FILING DATE: December 12, 1997
OR APPLICATION NUMBER: 60/069,696
OR FILING DATE: December 16, 1997
OR APPLICATION NUMBER: 60/069,696
OR APPLICATION NUMBER: 60/069,694
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FILING DATE: May 22, 2000
APPLICATION NUMBER: PCT/US00/20710
FILING DATE: July 28, 2000
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APPLICATION NUMBER: PCT/US00/05841
FILING DATE: Warch 2, 2000
APPLICATION NUMBER: PCT/US00/08439
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FILING DATE: MATCh 3, 1999
APPLICATION NUMBER: PCT/US99/12252
FILING DATE: June 22, 1999
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APPLICATION NUMBER: PCT/US00/04414
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APPLICATION NUMBER: 09/218,517
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: No. US20020142419Alember
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NUMBER: PCT/US99/28409
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                                               CURRENT APPLICATION NUMBER: US/09/943,762
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 09/866,028
PRIOR FILING DATE: 2001-05-25
PRIOR PRIOR APPLICATION NUMBER: 60/067,411
PRIOR FILING DATE: December 3, 1997
PRIOR APPLICATION NUMBER: 60/069,334
PRIOR FILING DATE: December 11, 1997
PRIOR APPLICATION NUMBER: 60/069335
PRIOR APPLICATION NUMBER: 60/069335
PRIOR APPLICATION NUMBER: 60/06937
PRIOR APPLICATION NUMBER: 60/069,278
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FILING DATE: December 11, 1997
APPLICATION NUMBER: 60/069,425
FILING DATE: December 12, 1997
APPLICATION NUMBER: 60/069,696
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APPLICATION
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Goddard, Audrey
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Tumas, Daniel
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Query Match
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OR APPLICATION NUMBER: 60/075,945
OR FILING DATE: February 25, 1998
OR APPLICATION NUMBER: 60/112,850
OR FILING DATE: December 16, 1998
OR APPLICATION NUMBER: 60/113,296
OR APPLICATION NUMBER: 60/113,296
OR FILING DATE: December 22, 1998
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FILING DATE: February
R OF SEQ ID NOS: 120
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FILING DATE: MATCH 3, 1999
APPLICATION NUMBER: PCT/US99/12252
FILING DATE: June 22, 1999
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FILING DATE: December 16, 1998
APPLICATION NUMBER: 09/218,517
FILING DATE: December 22, 1998
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FILING DATE
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APPLICATION NUMBER: PCT/US99/28409
FILING DATE: No. US20020142958Alember 30,
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         Conservative
                                                                                                                                                                                                                                                                   NUMBER: PC: March 30,
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NUMBER: PCT/US99/28301
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NUMBER: 60/074,
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NIMBER: PCT/US00/20710
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                                                                           ICQVEHPAVTGNLQTQRYLEVQYKPQVHIQMTYPLQGLTREGDAFELTCEAIGKPQPVMV 240
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